



T 5/58/B AN

SEQUENCE LISTING

<110> Gerlach, ~~Patricia L~~
MacDougall, John R
Smithson, Glennda
Millet, Isabelle
Stone, David
Gunther, Erik
Ellerman, Karen
Grosse, William M
Alsobrook II, John P
Lepley, Denise M
Burgess, Catherine E
Padigaru, Muralidhara
Kekuda, Ramesh
Spytek, Kimberly A
Leach, Martin D
Shimkets, Richard A

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Pro Ala Leu Met Lys Ile Lys His Val Ser Asn Phe Val Arg Lys Cys
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Phe Glu Val Arg Ser Leu Val Gly Cys Gly His Phe Ala Glu Val Gln
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Phe Tyr Leu Ser Lys Gln Leu Asp Glu Ala Ser Gly Ala Asn Asp Glu			
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1480

1485

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Leu Ser Gln Lys Gln Arg Ser Phe Val Thr Phe Arg Gly Glu Pro Ala
 35 40 45

Glu Gly Phe Asn His Leu Val Val Asp Glu Arg Thr Gly His Ile Tyr
 50 55 60

Leu Gly Ala Val Asn Arg Ile Tyr Lys Leu Ser Ser Asp Leu Lys Val
 65 70 75 80

Leu Val Thr His Glu Thr Gly Pro Asp Glu Asp Asn Pro Lys Cys Tyr
 85 90 95

Pro Pro Arg Ile Val Gln Thr Cys Asn Glu Pro Leu Thr Thr Asn
 100 105 110

Asn Val Asn Lys Met Leu Leu Ile Asp Tyr Lys Glu Asn Arg Leu Ile
 115 120 125

Ala Cys Gly Ser Leu Tyr Gln Gly Ile Cys Lys Leu Leu Arg Leu Glu
 130 135 140

Asp Leu Phe Lys Leu Gly Glu Pro Tyr His Lys Lys Glu His Tyr Leu
 145 150 155 160

Ser Gly Val Asn Glu Ser Gly Ser Val Phe Gly Val Ile Val Ser Tyr
 165 170 175

Ser Asn Leu Asp Asp Lys Leu Phe Ile Ala Thr Ala Val Asp Gly Lys
 180 185 190

Pro Glu Tyr Phe Pro Thr Ile Ser Ser Arg Lys Leu Thr Lys Asn Ser
 195 200 205

Glu Ala Asp Gly Met Phe Ala Tyr Val Phe His Asp Glu Phe Val Ala
 210 215 220

Ser Met Ile Lys Ile Pro Ser Asp Thr Phe Thr Ile Ile Pro Asp Phe
 225 230 235 240

Asp Ile Tyr Tyr Val Tyr Gly Phe Ser Ser Gly Asn Phe Val Tyr Phe
 245 250 255

Leu Thr Leu Gln Pro Glu Met Val Ser Pro Pro Gly Ser Thr Thr Lys
 260 265 270

Glu Gln Val Tyr Thr Ser Lys Leu Val Arg Leu Cys Lys Glu Asp Thr
 275 280 285

Ala Phe Asn Ser Tyr Val Glu Val Pro Ile Gly Cys Glu Arg Ser Gly
 290 295 300

Val Glu Tyr Arg Leu Leu Gln Ala Ala Tyr Leu Ser Lys Ala Gly Ala
 305 310 315 320

Val Leu Gly Arg Thr Leu Gly Val His Pro Asp Asp Asp Leu Leu Phe
 325 330 335

Thr Val Phe Ser Lys Gly Gln Lys Arg Lys Met Lys Ser Leu Asp Glu
 340 345 350

Ser Ala Leu Cys Ile Phe Ile Leu Lys Gln Ile Asn Asp Arg Ile Lys
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 Glu Arg Leu Gln Ser Cys Tyr Arg Gly Glu Gly Thr Leu Asp Leu Ala
 370 375 380
 Trp Leu Lys Val Lys Asp Ile Pro Cys Ser Ser Ala Leu Leu Thr Ile
 385 390 395 400
 Asp Asp Asn Phe Cys Gly Leu Asp Met Asn Ala Pro Leu Gly Val Ser
 405 410 415
 Asp Met Val Arg Gly Ile Pro Val Phe Thr Glu Asp Arg Asp Arg Met
 420 425 430
 Thr Ser Val Ile Ala Tyr Val Tyr Lys Asn His Ser Leu Ala Phe Val
 435 440 445
 Gly Thr Lys Ser Gly Lys Leu Lys Lys Ile Arg Val Asp Gly Pro Arg
 450 455 460
 Gly Asn Ala Leu Gln Tyr Glu Thr Val Gln Val Val Asp Pro Gly Pro
 465 470 475 480
 Val Leu Arg Asp Met Ala Phe Ser Lys Asp His Glu Gln Leu Tyr Ile
 485 490 495
 Met Ser Glu Arg Gln Leu Thr Arg Val Pro Val Glu Ser Cys Gly Gln
 500 505 510
 Tyr Gln Ser Cys Gly Glu Cys Leu Gly Ser Gly Asp Pro His Cys Gly
 515 520 525
 Trp Cys Val Leu His Asn Thr Cys Thr Arg Lys Glu Arg Cys Glu Arg
 530 535 540
 Ser Lys Glu Pro Arg Arg Phe Ala Ser Glu Met Lys Gln Cys Val Arg
 545 550 555 560
 Leu Thr Val His Pro Asn Asn Ile Ser Val Ser Gln Tyr Asn Val Leu
 565 570 575
 Leu Val Leu Glu Thr Tyr Asn Val Pro Glu Leu Ser Ala Gly Val Asn
 580 585 590
 Cys Thr Phe Glu Asp Leu Ser Glu Met Asp Gly Leu Val Val Gly Asn
 595 600 605
 Gln Ile Gln Cys Tyr Ser Pro Ala Ala Lys Glu Val Pro Arg Ile Ile
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 Thr Glu Asn Gly Asp His His Val Val Gln Leu Gln Leu Lys Ser Lys
 625 630 635 640
 Glu Thr Gly Met Thr Phe Ala Ser Thr Ser Phe Val Phe Tyr Asn Cys
 645 650 655

Ser Val His Asn Ser Cys Leu Ser Cys Val Glu Ser Pro Tyr Arg Cys
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 His Trp Cys Lys Tyr Arg His Val Cys Thr His Asp Pro Lys Thr Cys
 675 680 685
 Ser Phe Gln Glu Gly Arg Val Lys Leu Pro Glu Asp Cys Pro Gln Leu
 690 695 700
 Leu Arg Val Asp Lys Ile Leu Val Pro Val Glu Val Ile Lys Pro Ile
 705 710 715 720
 Thr Leu Lys Ala Lys Asn Leu Pro Gln Pro Gln Ser Gly Gln Arg Gly
 725 730 735
 Tyr Glu Cys Ile Leu Asn Ile Gln Gly Ser Glu Gln Arg Val Pro Ala
 740 745 750
 Leu Arg Phe Asn Ser Ser Val Gln Cys Gln Asn Thr Ser Tyr Ser
 755 760 765
 Tyr Glu Gly Met Glu Ile Asn Asn Leu Pro Val Glu Leu Thr Val Val
 770 775 780
 Trp Asn Gly His Phe Asn Ile Asp Asn Pro Ala Gln Asn Lys Val His
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 Leu Tyr Lys Cys Gly Ala Met Arg Glu Ser Cys Gly Leu Cys Leu Lys
 805 810 815
 Ala Asp Pro Asp Phe Ala Cys Gly Trp Cys Gln Gly Pro Gly Gln Cys
 820 825 830
 Thr Leu Arg Gln His Cys Pro Ala Gln Glu Ser Gln Trp Leu Glu Leu
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 Ser Gly Ala Lys Ser Lys Cys Thr Asn Pro Arg Ile Thr Glu Ile Ile
 850 855 860
 Pro Val Thr Gly Pro Arg Glu Gly Gly Thr Lys Val Thr Ile Arg Gly
 865 870 875 880
 Glu Asn Leu Gly Leu Glu Phe Arg Asp Ile Ala Ser His Val Lys Val
 885 890 895
 Ala Gly Val Glu Cys Ser Pro Leu Val Asp Gly Tyr Ile Pro Ala Glu
 900 905 910
 Gln Ile Val Cys Glu Met Gly Glu Ala Lys Pro Ser Gln His Ala Gly
 915 920 925
 Phe Val Glu Ile Cys Val Ala Val Cys Arg Pro Glu Phe Met Ala Arg
 930 935 940
 Ser Ser Gln Leu Tyr Tyr Phe Met Thr Leu Thr Leu Ser Asp Leu Lys
 945 950 955 960

Pro Ser Arg Gly Pro Met Ser Gly Gly Thr Gln Val Thr Ile Thr Gly
 965 970 975
 Thr Asn Leu Asn Ala Gly Ser Asn Val Val Val Met Phe Gly Lys Gln
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 Pro Cys Leu Phe His Arg Arg Ser Pro Ser Tyr Ile Val Cys Asn Thr
 995 1000 1005
 Thr Ser Ser Asp Glu Val Leu Glu Met Lys Val Ser Val Gln Val Asp
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 Arg Ala Lys Ile His Gln Asp Leu Val Phe Gln Tyr Val Glu Asp Pro
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 1045 1050 1055
 Pro Ile Ala Val Trp Gly Thr His Leu Asp Leu Ile Gln Asn Pro Gln
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 Ile Arg Ala Lys His Gly Gly Lys Glu His Ile Asn Ile Cys Glu Val
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 Pro Asp His Gln Ser Asp Leu Thr Glu Arg Pro Glu Glu Phe Gly Phe
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 Ile Leu Asp Asn Val Gln Ser Leu Leu Ile Leu Asn Lys Thr Asn Phe
 1125 1130 1135
 Thr Tyr Tyr Pro Asn Pro Val Phe Glu Ala Phe Gly Pro Ser Gly Ile
 1140 1145 1150
 Leu Glu Leu Lys Pro Gly Thr Pro Ile Ile Leu Lys Gly Lys Asn Leu
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 Ile Pro Pro Val Ala Gly Gly Asn Val Lys Leu Asn Tyr Thr Val Leu
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 Val Gly Glu Lys Pro Cys Thr Val Thr Val Ser Asp Val Gln Leu Leu
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 Cys Glu Ser Pro Asn Leu Ile Gly Arg His Lys Val Met Ala Arg Val
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 Gly Gly Met Glu Tyr Ser Pro Gly Met Val Tyr Ile Ala Pro Asp Ser
 1220 1225 1230
 Pro Leu Ser Leu Pro Ala Ile Val Ser Ile Ala Val Ala Gly Gly Leu
 1235 1240 1245
 Leu Ile Ile Phe Ile Val Ala Val Leu Ile Ala Tyr Lys Arg Lys Ser
 1250 1255 1260

Arg Glu Ser Asp Leu Thr Leu Lys Arg Leu Gln Met Gln Met Asp Asn
 1265 1270 1275 1280

 Leu Glu Ser Arg Val Ala Leu Glu Cys Lys Glu Ala Phe Ala Glu Leu
 1285 1290 1295

 Gln Thr Asp Ile His Glu Leu Thr Ser Asp Leu Asp Gly Ala Gly Ile
 1300 1305 1310

 Pro Phe Leu Asp Tyr Arg Thr Tyr Thr Met Arg Val Leu Phe Pro Gly
 1315 1320 1325

 Ile Glu Asp His Pro Val Leu Arg Asp Leu Glu Val Pro Gly Tyr Arg
 1330 1335 1340

 Gln Glu Arg Val Glu Lys Gly Leu Lys Leu Phe Ala Gln Leu Ile Asn
 1345 1350 1355 1360

 Asn Lys Val Phe Leu Leu Ser Phe Ile Arg Thr Leu Glu Ser Gln Arg
 1365 1370 1375

 Ser Phe Ser Met Arg Asp Arg Gly Asn Val Ala Ser Leu Ile Met Thr
 1380 1385 1390

 Val Leu Gln Ser Lys Leu Glu Tyr Ala Thr Asp Val Leu Lys Gln Leu
 1395 1400 1405

 Leu Ala Asp Leu Ile Asp Lys Asn Leu Glu Ser Lys Asn His Pro Lys
 1410 1415 1420

 Leu Leu Leu Arg Arg Thr Glu Ser Val Ala Glu Lys Met Leu Thr Asn
 1425 1430 1435 1440

 Trp Phe Thr Phe Leu Leu Tyr Lys Phe Leu Lys Glu Cys Ala Gly Glu
 1445 1450 1455

 Pro Leu Phe Ser Leu Phe Cys Ala Ile Lys Gln Gln Met Glu Lys Gly
 1460 1465 1470

 Pro Ile Asp Ala Ile Thr Gly Glu Ala Arg Tyr Ser Leu Ser Glu Asp
 1475 1480 1485

 Lys Leu Ile Arg Gln Gln Ile Asp Tyr Lys Thr Leu Val Leu Ser Cys
 1490 1495 1500

 Val Ser Pro Asp Asn Ala Asn Ser Pro Glu Val Pro Val Lys Ile Leu
 1505 1510 1515 1520

 Asn Cys Asp Thr Ile Thr Gln Val Lys Glu Lys Ile Leu Asp Ala Ile
 1525 1530 1535

 Phe Lys Asn Val Pro Cys Ser His Arg Pro Lys Ala Ala Asp Met Asp
 1540 1545 1550

 Leu Glu Trp Arg Gln Gly Ser Gly Ala Arg Met Ile Leu Gln Asp Glu
 1555 1560 1565

Asp Ile Thr Thr Lys Ile Glu Asn Asp Trp Lys Arg Leu Asn Thr Leu
 1570 1575 1580

Ala His Tyr Gln Val Pro Asp Gly Ser Val Val Ala Leu Val Ser Lys
 1585 1590 1595 1600

Gln Val Thr Ala Tyr Asn Ala Val Asn Asn Ser Thr Val Ser Arg Thr
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Ser Ala Ser Lys Tyr Glu Asn Met Ile Arg Tyr Thr Gly Ser Pro Asp
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Ser Leu Arg Ser Arg Thr Pro Met Ile Thr Pro Asp Leu Glu Ser Gly
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Val Lys Met Trp His Leu Val Lys Asn His Glu His Gly Asp Gln Lys
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Glu Gly Asp Arg Gly Ser Lys Met Val Ser Glu Ile Tyr Leu Thr Arg
 1665 1670 1675 1680

Leu Leu Ala Thr Lys Gly Thr Leu Gln Lys Phe Val Asp Asp Leu Phe
 1685 1690 1695

Glu Thr Ile Phe Ser Thr Ala His Arg Gly Ser Ala Leu Pro Leu Ala
 1700 1705 1710

Ile Lys Tyr Met Phe Asp Phe Leu Asp Glu Gln Ala Asp Lys His Gly
 1715 1720 1725

Ile His Asp Pro His Val Arg His Thr Trp Lys Ser Asn Cys Leu Pro
 1730 1735 1740

Leu Arg Phe Trp Val Asn Met Ile Lys Asn Pro Gln Phe Val Phe Asp
 1745 1750 1755 1760

Ile His Lys Asn Ser Ile Thr Asp Ala Cys Leu Ser Val Val Ala Gln
 1765 1770 1775

Thr Phe Met Asp Ser Cys Ser Thr Ser Glu His Arg Leu Gly Lys Asp
 1780 1785 1790

Ser Pro Ser Asn Lys Leu Leu Tyr Ala Lys Asp Ile Pro Ser Tyr Lys
 1795 1800 1805

Asn Trp Val Glu Arg Tyr Tyr Ser Asp Ile Gly Lys Met Pro Ala Ile
 1810 1815 1820

Ser Asp Gln Asp Met Asn Ala Tyr Leu Ala Glu Gln Ser Arg Met His
 1825 1830 1835 1840

Met Asn Glu Phe Asn Thr Met Ser Ala Leu Ser Glu Ile Phe Ser Tyr
 1845 1850 1855

Val Gly Lys Tyr Ser Glu Glu Ile Leu Gly Pro Leu Asp His Asp Asp
 1860 1865 1870

Gln Cys Gly Lys Gln Lys Leu Ala Tyr Lys Leu Glu Gln Val Ile Thr
1875 1880 1885

Leu Met Ser Leu Asp Ser Asn Lys
1890 1895

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<211> 1535
<212> DNA
<213> Homo sapiens

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<211> 480
<212> PRT
<213> Homo sapiens

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Ala Pro Pro Leu Gly Pro Val Gln Val Val Thr Ala Cys Leu Leu Thr
35 40 45

Leu	Leu	Val	Ile	Trp	Thr	Leu	Leu	Gly	Asn	Val	Leu	Val	Ser	Ala	Ala
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Ile	Val	Trp	Ser	Arg	His	Leu	Arg	Ala	Lys	Met	Thr	Asn	Val	Phe	Ile
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Val	Ser	Leu	Pro	Val	Ser	Asp	Leu	Phe	Val	Ala	Leu	Leu	Val	Met	Ser
					85				90				95		
Trp	Lys	Ala	Val	Ala	Glu	Val	Ala	Gly	Tyr	Trp	Pro	Phe	Glu	Ala	Phe
					100				105				110		
Cys	Asp	Val	Trp	Val	Ala	Phe	Asp	Ile	Met	Cys	Ser	Thr	Ala	Ser	Ile
					115				120				125		
Leu	Asn	Leu	Cys	Val	Ile	Ser	Val	Ala	Arg	Tyr	Trp	Ala	Ile	Ser	Arg
					130				135				140		
Pro	Phe	Arg	Tyr	Glu	Arg	Lys	Met	Thr	Gln	Arg	Met	Ala	Leu	Val	Met
					145				150				155		160
Val	Arg	Pro	Ala	Trp	Thr	Leu	Ser	Ser	Leu	Ile	Ser	Phe	Ile	Pro	Val
					165				170				175		
Gln	Leu	Asn	Trp	His	Arg	Asp	Gln	Ala	Val	Ser	Trp	Gly	Gly	Leu	Asp
					180				185				190		
Leu	Pro	Asn	Asn	Leu	Ala	Asn	Trp	Thr	Pro	Trp	Glu	Glu	Ala	Val	Trp
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Glu	Pro	Asp	Val	Arg	Ala	Glu	Asn	Cys	Asp	Ser	Ser	Leu	Asn	Arg	Thr
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Tyr	Ala	Ile	Pro	Ser	Ser	Leu	Ile	Ser	Phe	Tyr	Ile	Pro	Met	Ala	Ile
					225				230				235		240
Met	Ile	Val	Thr	Tyr	Thr	Arg	Ile	Tyr	Arg	Ile	Ala	Gln	Val	Gln	Ile
					245				250				255		
Arg	Arg	Ile	Ser	Ser	Leu	Glu	Arg	Ala	Ala	Glu	His	Val	Gln	Ser	Cys
					260				265				270		
Arg	Ser	Ser	Ala	Gly	Cys	Thr	Pro	Asp	Thr	Ser	Leu	Arg	Phe	Ser	Ile
					275				280				285		
Lys	Lys	Glu	Thr	Glu	Val	Leu	Lys	Thr	Leu	Ser	Val	Ile	Met	Gly	Val
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Phe	Val	Cys	Cys	Trp	Leu	Pro	Phe	Phe	Ile	Leu	Asn	Cys	Met	Val	Pro
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Phe	Cys	Ser	Gly	His	Pro	Lys	Gly	Pro	Pro	Ala	Gly	Phe	Pro	Cys	Val
					325				330				335		
Ser	Glu	Thr	Thr	Phe	Asp	Val	Phe	Ile	Trp	Phe	Cys	Trp	Ala	Asn	Ser
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Ser Leu Asn Pro Val Pro Ser His Tyr Ala Phe Asn Ala Asp Phe Arg
 355 360 365

 Lys Val Phe Ala Gln Leu Leu Gly Cys Ser His Val Cys Ser Arg Thr
 370 375 380

 Pro Val Glu Thr Val Asn Ile Ser Asn Glu Leu Ile Ser Tyr Asn Gln
 385 390 395 400

 Asp Thr Val Phe His Lys Glu Ile Ala Ala Ala Tyr Ile His Met Met
 405 410 415

 Pro Asn Ala Ile Pro Pro Gly Asp Arg Glu Val Asp Asn Asp Glu Glu
 420 425 430

 Glu Glu Ser Pro Phe Asp Arg Met Ser Gln Ile Tyr Gln Thr Ser Pro
 435 440 445

 Asp Gly Asp His Val Ala Glu Ser Val Trp Glu Leu Asp Cys Glu Gly
 450 455 460

 Glu Ile Ser Leu Asp Lys Ile Thr Pro Phe Thr Pro Asn Gly Phe His
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<210> 16
 <211> 2657
 <212> DNA
 <213> Homo sapiens

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<210> 17

<211> 879

<212> PRT

<213> Homo sapiens

<400> 17

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															30
20															

Ile	Lys	Ile	Glu	Gly	Asp	Leu	Val	Ley	Gly	Ley	Phe	Pro	Ile	Asn
35														45

Glu	Lys	Gly	Thr	Gly	Thr	Glu	Glu	Cys	Gly	Arg	Ile	Asn	Glu	Asp	Arg
50															60

Gly	Ile	Gln	Arg	Leu	Glu	Ala	Met	Leu	Phe	Ala	Ile	Asp	Glu	Ile	Asn
65															80

Lys	Asp	Asn	Tyr	Leu	Leu	Pro	Gly	Val	Lys	Leu	Gly	Val	His	Ile	Leu
85															95

Asp	Thr	Cys	Ser	Arg	Asp	Thr	Tyr	Ala	Leu	Glu	Gln	Ser	Leu	Glu	Phe
100															110

Val	Arg	Ala	Ser	Leu	Thr	Lys	Val	Asp	Glu	Ala	Glu	Tyr	Met	Cys	Pro
115															125

Asp	Gly	Ser	Tyr	Ala	Ile	Gln	Glu	Asn	Ile	Pro	Leu	Leu	Ile	Ala	Gly
130															140

Val Ile Gly Gly Ser Tyr Ser Ser Val Ser Ile Gln Val Ala Asn Leu
 145 150 155 160

Leu Arg Leu Phe Gln Ile Pro Gln Ile Ser Tyr Ala Ser Thr Ser Ala
 165 170 175

Lys Leu Ser Asp Lys Ser Arg Tyr Asp Tyr Phe Ala Arg Thr Val Pro
 180 185 190

Pro Asp Phe Tyr Gln Ala Lys Ala Met Ala Glu Ile Leu Arg Tyr Phe
 195 200 205

Asn Trp Thr Tyr Val Ser Thr Val Ala Ser Glu Gly Asp Tyr Gly Glu
 210 215 220

Thr Gly Ile Glu Ala Phe Glu Gln Glu Ala Arg Leu Arg Asn Ile Cys
 225 230 235 240

Ile Ala Thr Ala Glu Lys Val Gly Arg Ser Asn Ile Arg Lys Ser Tyr
 245 250 255

Asp Ser Val Ile Arg Glu Leu Leu Gln Lys Pro Asn Ala Arg Val Val
 260 265 270

Val Leu Phe Met Arg Ser Asp Asp Ser Arg Glu Leu Ile Ala Ala Ala
 275 280 285

Ser Arg Val Asn Ala Ser Phe Thr Trp Val Ala Ser Asp Gly Trp Gly
 290 295 300

Ala Gln Glu Ser Ile Val Lys Gly Ser Glu His Val Ala Tyr Gly Ala
 305 310 315 320

Ile Thr Leu Glu Leu Ala Ser His Pro Val Arg Gln Phe Asp Arg Tyr
 325 330 335

Phe Gln Ser Leu Asn Pro Tyr Asn Asn His Arg Asn Pro Trp Phe Arg
 340 345 350

Asp Phe Trp Glu Gln Lys Phe Gln Cys Ser Leu Gln Asn Lys Arg Asn
 355 360 365

His Arg Gln Ile Cys Asp Lys His Leu Ala Ile Asp Ser Ser Asn Tyr
 370 375 380

Glu Gln Glu Ser Lys Ile Met Phe Val Val Asn Ala Val Tyr Ala Met
 385 390 395 400

Ala His Ala Leu His Lys Met Gln Arg Thr Leu Cys Pro Asn Thr Thr
 405 410 415

Lys Leu Cys Asp Ala Met Lys Ile Leu Asp Gly Lys Lys Leu Tyr Lys
 420 425 430

Asp Tyr Leu Leu Lys Ile Asn Phe Leu Ala Pro Phe Asn Pro Asn Lys
 435 440 445

Gly Ala Asp Ser Ile Val Lys Phe Asp Thr Tyr Gly Asp Gly Met Gly
 450 455 460
 Arg Tyr Asn Val Phe Asn Phe Gln His Ile Gly Gly Lys Tyr Ser Tyr
 465 470 475 480
 Leu Lys Val Gly His Trp Ala Glu Thr Leu Tyr Leu Asp Val Asp Ser
 485 490 495
 Ile His Trp Ser Arg Asn Ser Val Pro Thr Ser Gln Cys Ser Asp Pro
 500 505 510
 Cys Ala Pro Asn Glu Met Lys Asn Met Gln Pro Gly Asp Val Cys Cys
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 Trp Ile Cys Ile Pro Cys Glu Pro Tyr Glu Tyr Leu Val Asp Glu Phe
 530 535 540
 Thr Cys Met Asp Cys Gly Pro Gly Gln Trp Pro Thr Ala Asp Leu Ser
 545 550 555 560
 Gly Cys Tyr Asn Leu Pro Glu Asp Tyr Ile Arg Trp Glu Asp Ala Trp
 565 570 575
 Ala Ile Gly Pro Val Thr Ile Ala Cys Leu Gly Phe Met Cys Thr Cys
 580 585 590
 Ile Val Ile Thr Val Phe Ile Lys His Asn Asn Thr Pro Leu Val Lys
 595 600 605
 Ala Ser Gly Arg Glu Leu Cys Tyr Ile Leu Leu Phe Gly Val Ser Leu
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 Ser Tyr Cys Met Thr Phe Phe Ile Ala Lys Pro Ser Pro Val Ile
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 Cys Ala Leu Arg Arg Leu Gly Leu Gly Thr Ser Phe Ala Ile Cys Tyr
 645 650 655
 Ser Ala Leu Leu Thr Lys Thr Asn Cys Ile Ala Arg Ile Phe Asp Gly
 660 665 670
 Val Lys Asn Gly Ala Gln Arg Pro Lys Phe Ile Ser Pro Ser Ser Gln
 675 680 685
 Val Phe Ile Cys Leu Gly Leu Ile Leu Val Gln Ile Val Met Val Ser
 690 695 700
 Val Trp Leu Ile Leu Glu Thr Pro Gly Thr Arg Arg Tyr Thr Leu Pro
 705 710 715 720
 Glu Lys Arg Glu Thr Val Ile Leu Lys Cys Asn Val Lys Asp Ser Ser
 725 730 735
 Met Leu Ile Ser Leu Thr Tyr Asp Val Val Leu Val Ile Leu Cys Thr
 740 745 750

Val Tyr Ala Phe Lys Thr Arg Lys Cys Pro Glu Asn Phe Asn Glu Ala
755 760 765

Lys Phe Ile Gly Phe Thr Met Tyr Thr Thr Cys Ile Ile Trp Leu Ala
770 775 780

Phe Leu Pro Ile Phe Tyr Val Thr Ser Ser Asp Tyr Arg Val Gln Thr
785 790 795 800

Thr Thr Met Cys Ile Ser Val Ser Leu Ser Gly Phe Val Val Leu Gly
805 810 815

Cys Leu Phe Ala Pro Lys Val His Ile Val Leu Phe Gln Pro Gln Lys
820 825 830

Asn Val Val Thr His Arg Leu His Leu Asn Arg Phe Ser Val Ser Gly
835 840 845

Thr Ala Thr Thr Tyr Ser Gln Ala Ser Ala Ser Thr Tyr Val Pro Thr
850 855 860

Val Cys Asn Gly Arg Glu Val Leu Asp Ser Thr Thr Ser Ser Leu
865 870 875

<210> 18

<211> 1366

<212> DNA

<213> Homo sapiens

<400> 18

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<210> 19

<211> 442
<212> PRT
<213> Homo sapiens

<400> 19

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Ser Ser Arg Gly Cys Trp Tyr Tyr Leu Arg Tyr Phe Phe Leu Phe Val
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Ser Leu Ile Gln Phe Leu Ile Ile Leu Gly Leu Val Leu Phe Met Val
35 40 45

Tyr Gly Asn Val His Val Ser Thr Glu Ser Asn Leu Gln Ala Thr Glu
50 55 60

Arg Arg Ala Glu Gly Leu Tyr Ser Gln Leu Leu Gly Leu Thr Ala Ser
65 70 75 80

Gln Ser Asn Leu Thr Lys Glu Leu Asn Phe Thr Thr Arg Ala Lys Asp
85 90 95

Ala Ile Met Gln Met Trp Leu Asn Ala Arg Arg Asp Leu Asp Arg Ile
100 105 110

Asn Ala Ser Phe Arg Gln Cys Gln Gly Asp Arg Val Ile Tyr Thr Asn
115 120 125

Asn Gln Arg Tyr Met Ala Ala Ile Ile Leu Ser Glu Lys Gln Cys Arg
130 135 140

Asp Gln Phe Lys Asp Met Asn Lys Ser Cys Asp Ala Leu Leu Phe Met
145 150 155 160

Leu Asn Gln Lys Val Lys Thr Leu Glu Val Glu Ile Ala Lys Glu Lys
165 170 175

Thr Ile Cys Thr Lys Asp Lys Glu Ser Val Leu Leu Asn Lys Arg Val
180 185 190

Ala Glu Glu Gln Leu Val Glu Cys Val Lys Thr Arg Glu Leu Gln His
195 200 205

Gln Glu Arg Gln Leu Ala Lys Glu Gln Leu Gln Lys Val Gln Ala Leu
210 215 220

Cys Leu Pro Leu Asp Lys Asp Lys Phe Glu Met Asp Leu Arg Asn Leu
225 230 235 240

Trp Arg Asp Ser Ile Ile Pro Arg Ser Leu Asp Asn Leu Gly Tyr Asn
245 250 255

Leu Tyr His Pro Leu Gly Ser Glu Leu Ala Ser Ile Arg Arg Ala Cys
260 265 270

Asp His Met Pro Ser Leu Met Ser Ser Lys Val Glu Gly Gln Cys Arg

275

280

285

Ser Leu Arg Ala Asp Ile Glu Arg Val Ala Arg Glu Asn Ser Asp Leu
290 295 300

Gln Arg Gln Lys Leu Glu Ala Gln Gln Gly Leu Arg Ala Ser Gln Glu
305 310 315 320

Ala Lys Gln Lys Val Glu Lys Glu Ala Gln Ala Arg Glu Ala Lys Leu
325 330 335

Gln Ala Glu Cys Ser Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala
340 345 350

Val Leu Arg Lys Glu Arg Asp Asn Leu Ala Lys Glu Leu Glu Glu Lys
355 360 365

Lys Arg Glu Ala Glu Gln Leu Arg Met Glu Leu Ala Ile Arg Asn Ser
370 375 380

Ala Leu Asp Thr Cys Ile Lys Thr Lys Ser Gln Pro Met Met Pro Val
385 390 395 400

Ser Arg Pro Met Gly Pro Val Pro Asn Pro Gln Pro Ile Asp Pro Ala
405 410 415

Ser Leu Glu Glu Phe Lys Arg Lys Ile Leu Glu Ser Gln Arg Pro Pro
420 425 430

Ala Gly Ile Pro Val Ala Pro Ser Ser Gly
435 440

<210> 20
<211> 1421
<212> DNA
<213> Homo sapiens

<400> 20

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a a c t c a g c c c t g g a c a c t g c a t c a a g a c c a a g t c g c a g c c g a t g a t g c c a g t g t c a a g g 1260
c c c a t g g g c c c t g t c c c c a a c c c a g g c c a t c g a c c a g g c t g g a g g a t t c a a g g 1320
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<210> 21
<211> 442
<212> PRT
<213> *Homo sapiens*

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20 25 30

Ser Leu Ile Gln Phe Leu Ile Ile Leu Gly Leu Val Leu Phe Met Val
35 40 45

Tyr Gly Asp Val His Val Ser Thr Glu Ser Asn Leu Gln Ala Thr Glu
50 55 60

Arg Arg Ala Glu Gly Leu Tyr Ser Gln Leu Leu Gly Leu Thr Ala Ser
65 70 75 80

Gln Ser Asn Leu Thr Lys Glu Leu Asn Phe Thr Thr Arg Ala Lys Asp
85 90 95

Ala Ile Met Gln Met Trp Leu Asn Ala Arg Arg Asp Leu Asp Arg Ile
100 105 110

Asn Ala Ser Phe Arg Gln Cys Gln Gly Asp Arg Val Ile Tyr Thr Asn
115 120 125

Asn Gln Arg Tyr Met Ala Ala Ile Ile Leu Ser Glu Lys Gln Cys Arg
130 135 140

Asp Gln Phe Lys Asp Met Asn Lys Ser Cys Asp Ala Leu Leu Phe Met
145 150 155 160

Leu Asn Gln Lys Val Lys Thr Leu Glu Val Glu Ile Ala Lys Glu Lys
165 170 175

Thr Ile Cys Thr Lys Asp Lys Glu Ser Val Leu Leu Asn Lys Arg Val
 180 185 190

Ala Glu Glu Gln Leu Val Glu Cys Val Lys Thr Arg Glu Leu Gln His
195 200 205

Gln Glu Arg Gln Leu Ala Lys Glu Gln Leu Gln Lys Val Gln Ala Leu
210 215 220

Cys Leu Pro Leu Asp Lys Asp Lys Phe Glu Met Asp Leu Arg Asp Leu

225	230	235	240
Trp Arg Asp Ser Ile Ile Pro Arg Ser Leu Asp Asn Leu Gly Tyr Asn			
245		250	255
Leu Tyr His Pro Leu Gly Ser Glu Leu Ala Ser Ile Arg Arg Ala Cys			
260		265	270
Asp His Met Pro Ser Leu Met Ser Ser Lys Val Glu Glu Leu Ala Arg			
275	280		285
Ser Leu Arg Ala Asp Ile Glu Arg Val Ala Arg Glu Asn Ser Asp Leu			
290	295		300
Gln Arg Gln Lys Leu Glu Ala Gln Gln Gly Leu Arg Ala Ser Gln Glu			
305	310	315	320
Ala Lys Gln Lys Val Glu Lys Glu Ala Gln Ala Arg Glu Ala Lys Leu			
325		330	335
Gln Ala Glu Cys Ser Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala			
340		345	350
Val Leu Arg Lys Glu Arg Asp Asn Leu Ala Lys Glu Leu Glu Glu Lys			
355	360		365
Lys Arg Glu Ala Glu Gln Leu Arg Met Glu Leu Ala Ile Arg Asn Ser			
370	375		380
Ala Leu Asp Thr Cys Ile Lys Thr Lys Ser Gln Pro Met Met Pro Val			
385	390	395	400
Ser Arg Pro Met Gly Pro Val Pro Asn Pro Gln Pro Ile Asp Pro Ala			
405		410	415
Ser Leu Glu Glu Phe Lys Arg Lys Ile Leu Glu Ser Gln Arg Pro Pro			
420		425	430
Ala Gly Ile Pro Val Ala Pro Ser Ser Gly			
435		440	

<210> 22
 <211> 2024
 <212> DNA
 <213> Homo sapiens

<400> 22
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<210> 23
 <211> 442
 <212> PRT
 <213> Homo sapiens

<400> 23

Met Gly Leu Ala Met Glu His Gly Ser Tyr Ala Arg Ala Gly Gly
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Ser Ser Arg Gly Cys Trp Tyr Tyr Leu Arg Tyr Phe Phe Leu Phe Val
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Ser Leu Ile Gln Phe Leu Ile Ile Leu Gly Leu Val Leu Phe Met Val
 35 40 45

Tyr Gly Asn Val His Val Ser Thr Glu Ser Asn Leu Gln Ala Thr Glu
 50 55 60

Arg Arg Ala Glu Gly Leu Tyr Ser Gln Leu Leu Gly Leu Thr Ala Ser
 65 70 75 80

Gln Ser Asn Leu Thr Lys Glu Leu Asn Phe Thr Thr Arg Ala Lys Asp
 85 90 95

Ala Ile Met Gln Met Trp Leu Asn Ala Arg Arg Asp Leu Asp Arg Ile
 100 105 110

Asn Ala Ser Phe Arg Gln Cys Gln Gly Asp Arg Val Ile Tyr Thr Asn
 115 120 125

Asn Gln Arg Tyr Met Ala Ala Ile Ile Leu Ser Glu Lys Gln Cys Arg
 130 135 140
 Asp Gln Phe Lys Asp Met Asn Lys Ser Cys Asp Ala Leu Leu Phe Met
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 Leu Asn Gln Lys Val Lys Thr Leu Glu Val Glu Ile Ala Lys Glu Lys
 165 170 175
 Thr Ile Cys Thr Lys Asp Lys Glu Ser Val Leu Leu Asn Lys Arg Val
 180 185 190
 Ala Glu Glu Gln Leu Val Glu Cys Val Lys Thr Arg Glu Leu Gln His
 195 200 205
 Gln Glu Arg Gln Leu Ala Lys Glu Gln Leu Gln Arg Val Gln Ala Leu
 210 215 220
 Cys Leu Pro Leu Asp Lys Asp Lys Phe Glu Met Asp Leu Arg Asn Leu
 225 230 235 240
 Trp Arg Asp Ser Ile Ile Pro Arg Ser Leu Asp Asn Leu Gly Tyr Asn
 245 250 255
 Leu Tyr His Pro Leu Gly Ser Glu Leu Ala Ser Ile Arg Arg Ala Cys
 260 265 270
 Asp His Met Pro Ser Leu Val Ser Ser Lys Val Glu Glu Leu Ala Arg
 275 280 285
 Ser Leu Arg Ala Asp Ile Glu Arg Val Ala Arg Glu Asn Ser Asp Leu
 290 295 300
 Gln Arg Gln Lys Leu Glu Ala Gln Gln Gly Leu Arg Ala Ser Gln Glu
 305 310 315 320
 Ala Lys Gln Lys Val Glu Lys Glu Ala Gln Ala Arg Glu Ala Lys Leu
 325 330 335
 Gln Ala Glu Cys Ser Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala
 340 345 350
 Val Leu Arg Lys Glu Arg Asp Asn Leu Ala Lys Glu Leu Glu Glu Lys
 355 360 365
 Lys Arg Glu Ala Glu Gln Leu Arg Met Glu Leu Ala Ile Arg Asn Ser
 370 375 380
 Ala Leu Asp Thr Cys Ile Lys Thr Lys Ser Gln Pro Met Met Pro Val
 385 390 395 400
 Ser Arg Pro Met Gly Pro Val Pro Asn Pro Gln Pro Ile Asp Pro Ala
 405 410 415
 Ser Leu Glu Glu Phe Lys Arg Lys Ile Leu Glu Ser Gln Arg Pro Pro
 420 425 430

Ala Gly Ile Pro Val Ala Pro Ser Ser Gly

435

440

<210> 24
<211> 8640
<212> DNA
<213> Homo sapiens

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Arg Leu Cys Gln Ala Ala Ile Gln Lys Leu Gln Glu Tyr Ile Gln Leu
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Glu	Thr	Val	Gly	Leu	Ser	Phe	Gly	Asn	Ile	Pro	Val	Phe	Gly	Asp	Tyr
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Gly	Glu	Lys	Arg	Arg	Gly	Gly	Lys	Arg	Lys	Thr	His	Gln	Gly	Pro	
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Val	Leu	Asp	Val	Gly	Cys	Ile	Trp	Val	Thr	Glu	Leu	Arg	Lys	Asn	Ser
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 Lys His Ser Leu Pro Gln Leu Leu Asp Ser Ser Ser Ala Ser Gln Glu
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															25
															30

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 Val Asn Ser Val Asn Val Arg His Ala Ala Leu Ser Lys Val His Ala
 755 760 765

 Ile Leu Ser Lys Cys Pro Pro Gly Pro Val Arg Leu Val Ile Gly Arg
 770 775 780

 His Pro Asn Pro Lys Val Ser Glu Gln Glu Met Asp Glu Val Ile Ala
 785 790 795 800

 Arg Ser Thr Tyr Gln Glu Ser Lys Glu Ala Asn Ser Ser Pro Gly Leu
 805 810 815

 Gly Thr Pro Leu Lys Ser Pro Ser Leu Ala Lys Lys Asp Ser Leu Ile
 820 825 830

 Ser Glu Ser Glu Leu Ser Gln Tyr Phe Ala His Asp Val Pro Gly Pro
 835 840 845

 Leu Ser Asp Phe Met Val Val Gly Ser Glu Asp Glu Asp His Pro Gly
 850 855 860

 Ser Gly Cys Ser Thr Ser Glu Glu Gly Ser Leu Pro Pro Ser Thr Ser
 865 870 875 880

 Thr His Lys Glu Pro Gly Lys Pro Arg Ala Asn Ser Leu Val Thr Leu
 885 890 895

 Gly Ser His Arg Ala Ser Gly Leu Phe His Lys Gln Val Thr Val Ala
 900 905 910

 Arg Gln Ala Ser Leu Pro Gly Ser Pro Gln Ala Leu Arg Asn Pro Leu
 915 920 925

 Leu Arg Gln Arg Lys Val Gly Cys Tyr Asp Ala Asn Asp Ala Ser Asp
 930 935 940

Glu Glu Glu Phe Asp Arg Glu Gly Asp Cys Ile Ser Leu Pro Gly Ala
 945 950 955 960

Leu Pro Gly Pro Ile Arg Pro Leu Ser Glu Asp Asp Pro Arg Arg Val
 965 970 975

Ser Ile Ser Ser Ser Lys Gly Met Asp Val His Asn Gln Glu Glu Arg
 980 985 990

Pro Arg Lys Thr Leu Val Ser Lys Ala Ile Ser Ala Pro Leu Leu Gly
 995 1000 1005

Ser Ser Val Asp Leu Glu Glu Ser Ile Pro Glu Gly Met Val Asp Ala
 1010 1015 1020

Ala Ser Tyr Ala Ala Asn Leu Thr Asp Ser Ala Glu Ala Pro Lys Gly
 1025 1030 1035 1040

Ser Pro Gly Ser Trp Trp Lys Lys Glu Leu Ser Gly Ser Ser Ala
 1045 1050 1055

Pro Lys Leu Glu Tyr Thr Val Arg Thr Asp Thr Gln Ser Pro Thr Asn
 1060 1065 1070

Thr Gly Ser Pro Ser Ser Pro Gln Gln Lys Ser Glu Gly Leu Gly Ser
 1075 1080 1085

Arg His Arg Pro Val Ala Arg Val Ser Pro His Cys Lys Arg Ser Glu
 1090 1095 1100

Ala Glu Ala Lys Pro Ser Gly Ser Gln Thr Val Asn Leu Thr Gly Arg
 1105 1110 1115 1120

Ala Asn Asp Pro Cys Asp Leu Asp Ser Arg Val Gln Ala Thr Ser Val
 1125 1130 1135

Lys Val Thr Val Ala Gly Phe Gln Pro Gly Gly Ala Val Glu Lys Glu
 1140 1145 1150

Ser Leu Gly Lys Leu Thr Thr Gly Asp Ala Cys Val Ser Thr Ser Cys
 1155 1160 1165

Glu Leu Ala Ser Ala Leu Ser His Leu Asp Ala Ser His Leu Thr Glu
 1170 1175 1180

Asn Leu Pro Lys Ala Ala Ser Glu Leu Gly Gln Gln Pro Met Thr Glu
 1185 1190 1195 1200

Leu Asp Ser Ser Ser Asp Leu Ile Ser Ser Pro Gly Lys Lys Gly Ala
 1205 1210 1215

Ala His Pro Asp Pro Ser Lys Thr Ser Val Asp Thr Gly Lys Val Ser
 1220 1225 1230

Arg Pro Glu Asn Pro Ser Gln Pro Ala Ser Pro Arg Val Ala Lys Cys
 1235 1240 1245

Lys Ala Arg Ser Pro Val Arg Leu Pro His Glu Gly Ser Pro Ser Pro
 1250 1255 1260

 Gly Glu Lys Ala Ala Ala Pro Pro Asp Tyr Ser Lys Thr Arg Ser Ala
 1265 1270 1275 1280

 Ser Glu Thr Ser Thr Pro His Asn Thr Arg Arg Val Ala Ala Leu Arg
 1285 1290 1295

 Gly Ala Gly Pro Gly Ala Glu Gly Met Thr Pro Ala Gly Ala Val Leu
 1300 1305 1310

 Pro Gly Asp Pro Leu Thr Ser Gln Glu Gln Arg Gln Gly Ala Pro Gly
 1315 1320 1325

 Asn His Ser Lys Ala Leu Glu Met Thr Gly Ile His Ala Pro Glu Ser
 1330 1335 1340

 Ser Gln Glu Pro Ser Leu Leu Glu Gly Ala Asp Ser Val Ser Ser Arg
 1345 1350 1355 1360

 Ala Pro Gln Ala Ser Leu Ser Met Leu Pro Ser Thr Asp Asn Thr Lys
 1365 1370 1375

 Glu Ala Cys Gly His Val Ser Gly His Cys Cys Pro Gly Gly Ser Arg
 1380 1385 1390

 Glu Ser Pro Val Thr Asp Ile Asp Ser Phe Ile Lys Glu Leu Asp Ala
 1395 1400 1405

 Ser Ala Ala Arg Ser Pro Ser Ser Gln Thr Gly Asp Ser Gly Ser Gln
 1410 1415 1420

 Glu Gly Ser Ala Gln Gly His Pro Pro Ala Gly Ala Gly Gly Ser
 1425 1430 1435 1440

 Ser Cys Arg Ala Glu Pro Val Pro Gly Gly Gln Thr Ser Ser Pro Arg
 1445 1450 1455

 Arg Ala Trp Ala Ala Gly Ala Pro Ala Tyr Pro Gln Trp Ala Ser Gln
 1460 1465 1470

 Pro Ser Val Leu Asp Ser Ile Asn Pro Asp Lys His Phe Thr Val Asn
 1475 1480 1485

 Lys Asn Phe Leu Ser Asn Tyr Ser Arg Asn Phe Ser Ser Phe His Glu
 1490 1495 1500

 Asp Ser Thr Ser Leu Ser Gly Leu Gly Asp Ser Thr Glu Pro Ser Leu
 1505 1510 1515 1520

 Ser Ser Met Tyr Gly Asp Ala Glu Asp Ser Ser Ser Asp Pro Glu Ser
 1525 1530 1535

 Leu Thr Glu Ala Pro Arg Ala Ser Ala Arg Asp Gly Trp Ser Pro Pro
 1540 1545 1550

Arg Ser Arg Val Ser Leu His Lys Glu Asp Pro Ser Glu Ser Glu Glu
 1555 1560 1565

Glu Gln Ile Glu Ile Cys Ser Thr Arg Gly Cys Pro Asn Pro Pro Ser
 1570 1575 1580

Ser Pro Ala His Leu Pro Thr Gln Ala Ala Ile Cys Pro Ala Ser Ala
 1585 1590 1595 1600

Lys Val Leu Ser Leu Lys Tyr Ser Thr Pro Arg Glu Ser Val Ala Ser
 1605 1610 1615

Pro Arg Glu Lys Val Ala Cys Leu Pro Gly Ser Tyr Thr Ser Gly Pro
 1620 1625 1630

Asp Ser Ser Gln Pro Ser Ser Leu Leu Glu Met Ser Ser Gln Glu His
 1635 1640 1645

Glu Thr His Ala Asp Ile Ser Thr Ser Gln Asn His Arg Pro Ser Cys
 1650 1655 1660

Ala Glu Glu Thr Thr Glu Val Thr Ser Ala Ser Ser Ala Met Glu Asn
 1665 1670 1675 1680

Ser Pro Leu Ser Lys Val Ala Arg His Phe His Ser Pro Pro Ile Ile
 1685 1690 1695

Leu Ser Ser Pro Asn Met Val Asn Gly Leu Glu His Asp Leu Leu Asp
 1700 1705 1710

Asp Glu Thr Leu Asn Gln Tyr Glu Thr Ser Ile Asn Ala Ala Ala Ser
 1715 1720 1725

Leu Ser Ser Phe Ser Val Asp Val Pro Lys Asn Gly Glu Ser Val Leu
 1730 1735 1740

Glu Asn Leu His Ile Ser Glu Ser Gln Asp Leu Asp Asp Leu Leu Gln
 1745 1750 1755 1760

Lys Pro Lys Met Ile Ala Arg Arg Pro Ile Met Ala Trp Phe Lys Glu
 1765 1770 1775

Ile Asn Lys His Asn Gln Gly Thr His Leu Arg Ser Lys Thr Glu Lys
 1780 1785 1790

Glu Gln Pro Leu Met Pro Ala Arg Ser Pro Asp Ser Lys Ile Gln Met
 1795 1800 1805

Val Ser Ser Ser Gln Lys Lys Gly Val Thr Val Pro His Ser Pro Pro
 1810 1815 1820

Gln Pro Lys Thr Asn Leu Glu Asn Lys Asp Leu Ser Lys Lys Ser Pro
 1825 1830 1835 1840

Ala Glu Met Leu Leu Thr Asn Gly Gln Lys Ala Lys Cys Gly Pro Lys
 1845 1850 1855

Leu Lys Arg Leu Ser Leu Lys Gly Lys Ala Lys Val Asn Ser Glu Ala
 1860 1865 1870

 Pro Ala Ala Asn Ala Val Lys Ala Gly Gly Thr Asp His Arg Lys Pro
 1875 1880 1885

 Leu Ile Ser Pro Gln Thr Ser His Lys Thr Leu Ser Lys Ala Val Ser
 1890 1895 1900

 Gln Arg Leu His Val Ala Asp His Glu Asp Pro Asp Arg Asn Thr Thr
 1905 1910 1915 1920

 Ala Ala Pro Arg Ser Pro Gln Cys Val Leu Glu Ser Lys Pro Pro Leu
 1925 1930 1935

 Ala Thr Ser Gly Pro Leu Lys Pro Ser Val Ser Asp Thr Ser Ile Arg
 1940 1945 1950

 Thr Phe Val Ser Pro Leu Thr Ser Pro Lys Pro Val Pro Glu Gln Gly
 1955 1960 1965

 Met Trp Ser Arg Phe His Met Ala Val Leu Ser Glu Pro Asp Arg Gly
 1970 1975 1980

 Cys Pro Thr Thr Pro Lys Ser Pro Lys Cys Arg Ala Glu Gly Arg Ala
 1985 1990 1995 2000

 Pro Arg Ala Asp Ser Gly Pro Val Ser Pro Ala Ala Ser Arg Asn Gly
 2005 2010 2015

 Met Ser Val Ala Gly Asn Arg Gln Ser Glu Pro Arg Leu Ala Ser His
 2020 2025 2030

 Val Ala Ala Asp Thr Ala Gln Pro Arg Pro Thr Gly Glu Lys Gly Gly
 2035 2040 2045

 Asn Ile Met Ala Ser Asp Arg Leu Glu Arg Thr Asn Gln Leu Lys Ile
 2050 2055 2060

 Val Glu Ile Ser Ala Glu Ala Val Ser Glu Thr Val Cys Gly Asn Lys
 2065 2070 2075 2080

 Pro Ala Glu Ser Asp Arg Arg Gly Gly Cys Leu Ala Gln Gly Asn Cys
 2085 2090 2095

 Gln Glu Lys Ser Glu Ile Arg Leu Tyr Arg Gln Val Ala Glu Ser Ser
 2100 2105 2110

 Thr Ser His Pro Ser Ser Leu Pro Ser His Ala Ser Gln Ala Glu Gln
 2115 2120 2125

 Glu Met Ser Arg Ser Phe Ser Met Ala Lys Leu Ala Ser Ser Ser Ser
 2130 2135 2140

 Ser Leu Gln Thr Ala Ile Arg Lys Ala Glu Tyr Ser Gln Gly Lys Ser
 2145 2150 2155 2160

Ser Leu Met Ser Asp Ser Arg Gly Val Pro Arg Asn Ser Ile Pro Gly
 2165 2170 2175

 Gly Pro Ser Gly Glu Asp His Leu Tyr Phe Thr Pro Arg Pro Ala Thr
 2180 2185 2190

 Arg Thr Tyr Ser Met Pro Ala Gln Phe Ser Ser His Phe Gly Arg Glu
 2195 2200 2205

 Gly His Pro Pro His Ser Leu Gly Arg Ser Arg Asp Ser Gln Val Pro
 2210 2215 2220

 Val Thr Ser Ser Val Val Pro Glu Ala Lys Ala Ser Arg Gly Gly Leu
 2225 2230 2235 2240

 Pro Ser Leu Ala Asn Gly Gln Gly Ile Tyr Ser Val Lys Pro Leu Leu
 2245 2250 2255

 Asp Thr Ser Arg Asn Leu Pro Ala Thr Asp Glu Gly Asp Ile Ile Ser
 2260 2265 2270

 Val Gln Glu Thr Ser Cys Leu Val Thr Asp Lys Ile Lys Val Thr Arg
 2275 2280 2285

 Arg His Tyr Cys Tyr Glu Gln Asn Trp Pro His Glu Ser Thr Ser Phe
 2290 2295 2300

 Phe Ser Val Lys Gln Arg Ile Lys Ser Phe Glu Asn Leu Ala Asn Ala
 2305 2310 2315 2320

 Asp Arg Pro Val Ala Lys Ser Gly Ala Ser Pro Phe Leu Ser Val Ser
 2325 2330 2335

 Ser Lys Pro Pro Ile Gly Arg Arg Ser Ser Gly Ser Ile Val Ser Gly
 2340 2345 2350

 Ser Leu Gly His Pro Gly Asp Ala Ala Ala Arg Leu Leu Arg Arg Ser
 2355 2360 2365

 Leu Ser Ser Cys Ser Glu Asn Gln Ser Glu Ala Gly Thr Leu Leu Pro
 2370 2375 2380

 Gln Met Ala Lys Ser Pro Ser Ile Met Thr Leu Thr Ile Ser Arg Gln
 2385 2390 2395 2400

 Asn Pro Pro Glu Thr Ser Ser Lys Gly Ser Asp Ser Glu Leu Lys Lys
 2405 2410 2415

 Ser Leu Gly Pro Leu Gly Ile Pro Thr Pro Thr Met Thr Leu Ala Ser
 2420 2425 2430

 Pro Val Lys Arg Asn Lys Ser Ser Val Arg His Thr Gln Pro Ser Pro
 2435 2440 2445

 Val Ser Arg Ser Lys Leu Gln Glu Leu Arg Ala Leu Ser Met Pro Asp
 2450 2455 2460

Leu Asp Lys Leu Cys Ser Glu Asp Tyr Ser Ala Gly Pro Ser Ala Val
 2465 2470 2475 2480

 Leu Phe Lys Thr Glu Leu Glu Ile Thr Pro Arg Arg Ser Pro Gly Pro
 2485 2490 2495

 Pro Ala Gly Gly Val Ser Cys Pro Glu Lys Gly Gly Asn Arg Ala Cys
 2500 2505 2510

 Pro Gly Gly Ser Gly Pro Lys Thr Ser Ala Ala Glu Thr Pro Ser Ser
 2515 2520 2525

 Ala Ser Asp Thr Gly Glu Ala Ala Gln Asp Leu Pro Phe Arg Arg Ser
 2530 2535 2540

 Trp Ser Val Asn Leu Asp Gln Leu Leu Val Ser Ala Gly Asp Gln Gln
 2545 2550 2555 2560

 Arg Leu Gln Ser Val Leu Ser Ser Val Gly Ser Lys Ser Thr Ile Leu
 2565 2570 2575

 Thr Leu Ile Gln Glu Ala Lys Ala Gln Ser Glu Asn Glu Glu Asp Val
 2580 2585 2590

 Cys Phe Ile Val Leu Asn Arg Lys Glu Gly Ser Gly Leu Gly Phe Ser
 2595 2600 2605

 Val Ala Gly Gly Thr Asp Val Glu Pro Lys Ser Ile Thr Val His Arg
 2610 2615 2620

 Val Phe Ser Gln Gly Ala Ala Ser Gln Glu Gly Thr Met Asn Arg Gly
 2625 2630 2635 2640

 Asp Phe Leu Leu Ser Val Asn Gly Ala Ser Leu Ala Gly Leu Ala His
 2645 2650 2655

 Gly Asn Val Leu Lys Val Leu His Gln Ala Gln Leu His Lys Asp Ala
 2660 2665 2670

 Leu Val Val Ile Lys Lys Gly Met Asp Gln Pro Arg Pro Ser Ala Arg
 2675 2680 2685

 Gln Glu Pro Pro Thr Ala Asn Gly Lys Gly Leu Leu Ser Arg Lys Thr
 2690 2695 2700

 Ile Pro Leu Glu Pro Gly Ile Gly Arg Ser Val Ala Val His Asp Ala
 2705 2710 2715 2720

 Leu Cys Val Glu Val Leu Lys Thr Ser Ala Gly Leu Gly Leu Ser Leu
 2725 2730 2735

 Asp Gly Gly Lys Ser Ser Val Thr Gly Asp Gly Pro Leu Val Ile Lys
 2740 2745 2750

 Arg Val Tyr Lys Gly Gly Ala Ala Glu Gln Ala Gly Ile Ile Glu Ala
 2755 2760 2765

Gly Asp Glu Ile Leu Ala Ile Asn Gly Lys Pro Leu Val Gly Leu Met
2770 2775 2780

His Phe Asp Ala Trp Asn Ile Met Lys Ser Val Pro Glu Gly Pro Val
2785 2790 2795 2800

Gln Leu Leu Ile Arg Lys His Arg Asn Ser Ser
2805 2810

<210> 28
<211> 798
<212> PRT
<213> Homo sapiens

<400> 28
Met Leu Ile Asn Cys Glu Ala Lys Gly Ile Lys Met Val Ser Glu Ile
1 5 10 15

Ser Val Pro Pro Ser Arg Pro Phe Gln Leu Ser Leu Leu Asn Asn Gly
20 25 30

Leu Thr Met Leu His Thr Asn Asp Phe Ser Gly Leu Thr Asn Ala Ile
35 40 45

Ser Ile His Leu Gly Phe Asn Asn Ile Ala Asp Ile Glu Ile Gly Ala
50 55 60

Phe Asn Gly Leu Gly Leu Leu Lys Gln Leu His Ile Asn His Asn Ser
65 70 75 80

Leu Glu Ile Leu Lys Glu Asp Thr Phe His Gly Leu Glu Asn Leu Glu
85 90 95

Phe Leu Gln Ala Asp Asn Asn Phe Ile Thr Val Ile Glu Pro Ser Ala
100 105 110

Phe Ser Lys Leu Asn Arg Leu Lys Val Leu Ile Leu Asn Asp Asn Ala
115 120 125

Ile Glu Ser Leu Pro Pro Asn Ile Phe Arg Phe Val Pro Leu Thr His
130 135 140

Leu Asp Leu Arg Gly Asn Gln Leu Gln Thr Leu Pro Tyr Val Gly Phe
145 150 155 160

Leu Glu His Ile Gly Arg Ile Leu Asp Leu Gln Leu Glu Asp Asn Lys
165 170 175

Trp Ala Cys Asn Cys Asp Leu Leu Gln Leu Lys Thr Trp Leu Glu Asn
180 185 190

Met Pro Pro Gln Ser Ile Ile Gly Asp Val Val Cys Asn Ser Pro Pro
195 200 205

Phe Phe Lys Gly Ser Ile Leu Ser Arg Leu Lys Lys Glu Ser Ile Cys
210 215 220

Pro Thr Pro Pro Val Tyr Glu Glu His Glu Asp Pro Ser Gly Ser Leu
 225 230 235 240
 His Leu Ala Ala Thr Ser Ser Ile Asn Asp Ser Arg Met Ser Thr Lys
 245 250 255
 Thr Thr Ser Ile Leu Lys Leu Pro Thr Lys Ala Pro Gly Leu Ile Pro
 260 265 270
 Tyr Ile Thr Lys Pro Ser Thr Gln Leu Pro Gly Pro Tyr Cys Pro Ile
 275 280 285
 Pro Cys Asn Cys Lys Val Leu Ser Pro Ser Gly Leu Leu Ile His Cys
 290 295 300
 Gln Glu Arg Asn Ile Glu Ser Leu Ser Asp Leu Arg Pro Pro Pro Gln
 305 310 315 320
 Asn Pro Arg Lys Leu Ile Leu Ala Gly Asn Ile Ile His Ser Leu Met
 325 330 335
 Lys Ser Asp Leu Val Glu Tyr Phe Thr Leu Glu Met Leu His Leu Gly
 340 345 350
 Asn Asn Arg Ile Glu Val Leu Glu Glu Gly Ser Phe Met Asn Leu Thr
 355 360 365
 Arg Leu Gln Lys Leu Tyr Leu Asn Gly Asn His Leu Thr Lys Leu Ser
 370 375 380
 Lys Gly Met Phe Leu Gly Leu His Asn Leu Glu Tyr Leu Tyr Leu Glu
 385 390 395 400
 Tyr Asn Ala Ile Lys Glu Ile Leu Pro Gly Thr Phe Asn Pro Met Pro
 405 410 415
 Lys Leu Lys Val Leu Tyr Leu Asn Asn Asn Leu Leu Gln Val Leu Pro
 420 425 430
 Pro His Ile Phe Ser Gly Val Pro Leu Thr Lys Val Asn Leu Lys Thr
 435 440 445
 Asn Gln Phe Thr His Leu Pro Val Ser Asn Ile Leu Asp Asp Leu Asp
 450 455 460
 Leu Leu Thr Gln Ile Asp Leu Glu Asp Asn Pro Trp Asp Cys Ser Cys
 465 470 475 480
 Asp Leu Val Gly Leu Gln Gln Trp Ile Gln Lys Leu Ser Lys Asn Thr
 485 490 495
 Val Thr Asp Asp Ile Leu Cys Thr Ser Pro Gly His Leu Asp Lys Lys
 500 505 510
 Glu Leu Lys Ala Leu Asn Ser Glu Ile Leu Cys Pro Gly Leu Val Asn
 515 520 525

Asn	Pro	Ser	Met	Pro	Thr	Gln	Thr	Ser	Tyr	Leu	Met	Val	Thr	Thr	Pro
530						535					540				
Ala	Thr	Thr	Thr	Asn	Thr	Ala	Asp	Thr	Ile	Leu	Arg	Ser	Leu	Thr	Asp
545						550				555					560
Ala	Val	Pro	Leu	Ser	Val	Leu	Ile	Leu	Gly	Leu	Leu	Ile	Met	Phe	Ile
							565			570					575
Thr	Ile	Val	Phe	Cys	Ala	Ala	Gly	Ile	Val	Val	Leu	Val	Leu	His	Arg
							580		585					590	
Arg	Arg	Arg	Tyr	Lys	Lys	Lys	Gln	Val	Asp	Glu	Gln	Met	Arg	Asp	Asn
							595		600				605		
Ser	Pro	Val	His	Leu	Gln	Tyr	Ser	Met	Tyr	Gly	His	Lys	Thr	Thr	His
							610		615			620			
His	Thr	Thr	Glu	Arg	Pro	Ser	Ala	Ser	Leu	Tyr	Glu	Gln	His	Met	Val
							625		630		635				640
Ser	Pro	Met	Val	His	Val	Tyr	Arg	Ser	Pro	Ser	Phe	Gly	Pro	Lys	His
							645		650				655		
Leu	Glu	Glu	Glu	Glu	Arg	Asn	Glu	Lys	Glu	Gly	Ser	Asp	Ala	Lys	
							660		665			670			
His	Leu	Gln	Arg	Ser	Leu	Leu	Glu	Gln	Glu	Asn	His	Ser	Pro	Leu	Thr
							675		680			685			
Gly	Ser	Asn	Met	Lys	Tyr	Lys	Thr	Thr	Asn	Gln	Ser	Thr	Glu	Phe	Leu
							690		695			700			
Ser	Phe	Gln	Asp	Ala	Ser	Ser	Leu	Tyr	Arg	Asn	Ile	Leu	Glu	Lys	Glu
							705		710		715				720
Arg	Glu	Leu	Gln	Gln	Leu	Gly	Ile	Thr	Glu	Tyr	Leu	Arg	Lys	Asn	Ile
							725		730			735			
Ala	Gln	Leu	Gln	Pro	Asp	Met	Glu	Ala	His	Tyr	Pro	Gly	Ala	His	Glu
							740		745			750			
Glu	Leu	Lys	Leu	Met	Glu	Thr	Leu	Met	Tyr	Ser	Arg	Pro	Arg	Lys	Val
							755		760			765			
Leu	Val	Glu	Gln	Thr	Lys	Asn	Glu	Tyr	Phe	Glu	Leu	Lys	Ala	Asn	Leu
							770		775			780			
His	Ala	Glu	Pro	Asp	Tyr	Leu	Glu	Val	Leu	Glu	Gln	Gln	Thr		
							785		790			795			

<210> 29
<211> 440
<212> PRT
<213> Homo sapiens

<400> 29
 Met Leu Ile Asn Cys Glu Ala Lys Gly Ile Lys Met Val Ser Glu Ile
 1 5 10 15
 Ser Val Pro Pro Ser Arg Pro Phe Gln Leu Ser Leu Leu Asn Asn Gly
 20 25 30
 Leu Thr Met Leu His Thr Asn Asp Phe Ser Gly Leu Thr Asn Ala Ile
 35 40 45
 Ser Ile His Leu Gly Phe Asn Asn Ile Ala Asp Ile Glu Ile Gly Ala
 50 55 60
 Phe Asn Gly Leu Gly Leu Leu Lys Gln Leu His Ile Asn His Asn Ser
 65 70 75 80
 Leu Glu Ile Leu Lys Glu Asp Thr Phe His Gly Leu Glu Asn Leu Glu
 85 90 95
 Phe Leu Gln Ala Asp Asn Asn Phe Ile Thr Val Ile Glu Pro Ser Ala
 100 105 110
 Phe Ser Lys Leu Asn Arg Leu Lys Val Leu Ile Leu Asn Asp Asn Ala
 115 120 125
 Ile Glu Ser Leu Pro Pro Asn Ile Phe Arg Phe Val Pro Leu Thr His
 130 135 140
 Leu Asp Leu Arg Gly Asn Gln Leu Gln Thr Leu Pro Tyr Val Gly Phe
 145 150 155 160
 Leu Glu His Ile Gly Arg Ile Leu Asp Leu Gln Leu Glu Asp Asn Lys
 165 170 175
 Trp Ala Cys Asn Cys Asp Leu Leu Gln Leu Lys Thr Trp Leu Glu Asn
 180 185 190
 Met Pro Pro Gln Ser Ile Ile Gly Asp Val Val Cys Asn Ser Pro Pro
 195 200 205
 Phe Phe Lys Gly Ser Ile Leu Ser Arg Leu Lys Lys Glu Ser Ile Cys
 210 215 220
 Pro Thr Pro Pro Val Tyr Glu Glu His Glu Asp Pro Ser Gly Ser Leu
 225 230 235 240
 His Leu Ala Ala Thr Ser Ser Ile Asn Asp Ser Arg Met Ser Thr Lys
 245 250 255
 Thr Thr Ser Ile Leu Lys Leu Pro Thr Lys Ala Pro Gly Leu Ile Pro
 260 265 270
 Tyr Ile Thr Lys Pro Ser Thr Gln Leu Pro Gly Pro Tyr Cys Pro Ile
 275 280 285
 Pro Cys Asn Cys Lys Val Leu Ser Pro Ser Gly Leu Leu Ile His Cys

290

295

300

Gln Glu Arg Asn Ile Glu Ser Leu Ser Asp Leu Arg Pro Pro Pro Gln
 305 310 315 320

Asn Pro Arg Lys Leu Ile Leu Ala Gly Asn Ile Ile His Ser Leu Met
 325 330 335

Lys Ser Asp Leu Val Glu Tyr Phe Thr Leu Glu Met Leu His Leu Gly
 340 345 350

Asn Asn Arg Ile Glu Val Leu Glu Glu Gly Ser Phe Met Asn Leu Thr
 355 360 365

Arg Leu Gln Lys Leu Tyr Leu Asn Gly Asn His Leu Thr Lys Leu Ser
 370 375 380

Lys Gly Met Phe Leu Gly Leu His Asn Leu Glu Tyr Leu Tyr Leu Glu
 385 390 395 400

Tyr Asn Ala Ile Lys Glu Ile Leu Pro Gly Thr Phe Asn Pro Met Pro
 405 410 415

Lys Leu Lys Val Leu Tyr Leu Asn Asn Thr Ser Ser Lys Phe Tyr His
 420 425 430

His Ile Phe Phe Gln Gly Phe Leu
 435 440

<210> 30

<211> 853

<212> PRT

<213> Homo sapiens

<400> 30

Tyr Phe Ser Leu Phe Arg Ser Ile Gln Leu Phe Ala Asp Cys Lys Lys
 1 5 10 15

Met Phe Leu Trp Leu Phe Leu Ile Leu Ser Ala Leu Ile Ser Ser Thr
 20 25 30

Asn Ala Asp Ser Asp Ile Ser Val Glu Ile Cys Asn Val Cys Ser Cys
 35 40 45

Val Ser Val Glu Asn Val Leu Tyr Val Asn Cys Glu Lys Val Ser Val
 50 55 60

Tyr Arg Pro Asn Gln Leu Lys Pro Pro Trp Ser Asn Phe Tyr His Leu
 65 70 75 80

Asn Phe Gln Asn Asn Phe Leu Asn Ile Leu Tyr Pro Asn Thr Phe Leu
 85 90 95

Asn Phe Ser His Ala Val Ser Leu His Leu Gly Asn Asn Lys Leu Gln
 100 105 110

Asn	Ile	Glu	Gly	Gly	Ala	Phe	Leu	Gly	Leu	Ser	Ala	Leu	Lys	Gln	Leu
		115				120						125			
His	Leu	Asn	Asn	Asn	Glu	Leu	Lys	Ile	Leu	Arg	Ala	Asp	Thr	Phe	Leu
		130				135				140					
Gly	Ile	Glu	Asn	Leu	Glu	Tyr	Leu	Gln	Ala	Asp	Tyr	Asn	Leu	Ile	Lys
		145				150				155				160	
Tyr	Ile	Glu	Arg	Gly	Ala	Phe	Asn	Lys	Leu	His	Lys	Leu	Lys	Val	Leu
		165				170				175					
Ile	Leu	Asn	Asp	Asn	Leu	Ile	Ser	Phe	Leu	Pro	Asp	Asn	Ile	Phe	Arg
		180				185				190					
Phe	Ala	Ser	Leu	Thr	His	Leu	Asp	Ile	Arg	Gly	Asn	Arg	Ile	Gln	Lys
		195				200				205					
Leu	Pro	Tyr	Ile	Gly	Val	Leu	Glu	His	Ile	Gly	Arg	Val	Val	Glu	Leu
		210				215				220					
Gln	Leu	Glu	Asp	Asn	Pro	Trp	Asn	Cys	Ser	Cys	Asp	Leu	Leu	Pro	Leu
		225				230				235			240		
Lys	Ala	Trp	Leu	Glu	Asn	Met	Pro	Tyr	Asn	Ile	Tyr	Ile	Gly	Glu	Ala
		245				250				255					
Ile	Cys	Glu	Thr	Pro	Ser	Asp	Leu	Tyr	Gly	Arg	Leu	Leu	Lys	Glu	Thr
		260				265				270					
Asn	Lys	Gln	Glu	Leu	Cys	Pro	Met	Gly	Thr	Gly	Ser	Asp	Phe	Asp	Val
		275				280				285					
Arg	Ile	Leu	Pro	Pro	Ser	Gln	Leu	Glu	Asn	Gly	Tyr	Thr	Thr	Pro	Asn
		290				295				300					
Gly	His	Thr	Thr	Gln	Thr	Ser	Leu	His	Arg	Leu	Val	Thr	Lys	Pro	Pro
		305				310				315			320		
Lys	Thr	Thr	Asn	Pro	Ser	Lys	Ile	Ser	Gly	Ile	Val	Ala	Gly	Lys	Ala
		325				330				335					
Leu	Ser	Asn	Arg	Asn	Leu	Ser	Gln	Ile	Val	Ser	Tyr	Gln	Thr	Arg	Val
		340				345				350					
Pro	Pro	Leu	Thr	Pro	Cys	Pro	Ala	Pro	Cys	Phe	Cys	Lys	Thr	His	Pro
		355				360				365					
Ser	Asp	Leu	Gly	Leu	Ser	Val	Asn	Cys	Gln	Glu	Lys	Asn	Ile	Gln	Ser
		370				375				380					
Met	Ser	Glu	Leu	Ile	Pro	Lys	Pro	Leu	Asn	Ala	Lys	Lys	Leu	His	Val
		385				390				395			400		
Asn	Gly	Asn	Ser	Ile	Lys	Asp	Val	Asp	Val	Ser	Asp	Phe	Thr	Asp	Phe
		405				410				415					

Glu Gly Leu Asp Leu Leu His Leu Gly Ser Asn Gln Ile Thr Val Ile
 420 425 430

 Lys Gly Asp Val Phe His Asn Leu Thr Asn Leu Arg Arg Leu Tyr Leu
 435 440 445

 Asn Gly Asn Gln Ile Glu Arg Leu Tyr Pro Glu Ile Phe Ser Gly Leu
 450 455 460

 His Asn Leu Gln Tyr Leu Tyr Leu Glu Tyr Asn Leu Ile Lys Glu Ile
 465 470 475 480

 Ser Ala Gly Thr Phe Asp Ser Met Pro Asn Leu Gln Leu Leu Tyr Leu
 485 490 495

 Asn Asn Asn Leu Leu Lys Ser Leu Pro Val Tyr Ile Phe Ser Gly Ala
 500 505 510

 Pro Leu Ala Arg Leu Asn Leu Arg Asn Asn Lys Phe Met Tyr Leu Pro
 515 520 525

 Val Ser Gly Val Leu Asp Gln Leu Gln Ser Leu Thr Gln Ile Asp Leu
 530 535 540

 Glu Gly Asn Pro Trp Asp Cys Thr Cys Asp Leu Val Ala Leu Lys Leu
 545 550 555 560

 Trp Val Glu Lys Leu Ser Asp Gly Ile Val Val Lys Glu Leu Lys Cys
 565 570 575

 Glu Thr Pro Val Gln Phe Ala Asn Ile Glu Leu Lys Ser Leu Lys Asn
 580 585 590

 Glu Ile Leu Cys Pro Lys Leu Leu Asn Lys Pro Ser Ala Pro Phe Thr
 595 600 605

 Ser Pro Ala Pro Ala Ile Thr Phe Thr Thr Pro Leu Gly Pro Ile Arg
 610 615 620

 Ser Pro Pro Gly Gly Pro Val Pro Leu Ser Ile Leu Ile Leu Ser Ile
 625 630 635 640

 Leu Val Val Leu Ile Leu Thr Val Phe Val Ala Phe Cys Leu Leu Val
 645 650 655

 Phe Val Leu Arg Arg Asn Lys Lys Pro Thr Val Lys His Glu Gly Leu
 660 665 670

 Gly Asn Pro Asp Cys Gly Ser Met Gln Leu Gln Leu Arg Lys His Asp
 675 680 685

 His Lys Thr Asn Lys Lys Asp Gly Leu Ser Thr Glu Ala Phe Ile Pro
 690 695 700

 Gln Thr Ile Glu Gln Met Ser Lys Ser His Thr Cys Gly Leu Lys Glu
 705 710 715 720

Ser Glu Thr Gly Phe Met Phe Ser Asp Pro Pro Gly Gln Lys Val Val
 725 730 735

 Met Arg Asn Val Ala Asp Lys Glu Lys Asp Leu Leu His Val Asp Thr
 740 745 750

 Arg Lys Arg Leu Ser Thr Ile Asp Glu Leu Asp Glu Leu Phe Pro Ser
 755 760 765

 Arg Asp Ser Asn Val Phe Ile Gln Asn Phe Leu Glu Ser Lys Lys Glu
 770 775 780

 Tyr Asn Ser Ile Gly Val Ser Gly Phe Glu Ile Arg Tyr Pro Glu Lys
 785 790 795 800

 Gln Pro Asp Lys Lys Ser Lys Lys Ser Leu Ile Gly Gly Asn His Ser
 805 810 815

 Lys Ile Val Val Glu Gln Arg Lys Ser Glu Tyr Phe Glu Leu Lys Ala
 820 825 830

 Lys Leu Gln Ser Ser Pro Asp Tyr Leu Gln Val Leu Glu Glu Gln Thr
 835 840 845

 Ala Leu Asn Lys Ile
 850

<210> 31
 <211> 314
 <212> PRT
 <213> Homo sapiens

<400> 31
 Leu Gln Gln Trp Ile Gln Lys Leu Ser Lys Asn Thr Val Thr Asp Asp
 1 5 10 15

 Ile Leu Cys Thr Ser Pro Gly His Leu Asp Lys Lys Glu Leu Lys Ala
 20 25 30

 Leu Asn Ser Glu Ile Leu Cys Pro Gly Leu Val Asn Asn Pro Ser Met
 35 40 45

 Pro Thr Gln Thr Ser Tyr Leu Met Val Thr Thr Pro Ala Thr Thr Thr
 50 55 60

 Asn Thr Ala Asp Thr Ile Leu Arg Ser Leu Thr Asp Ala Val Pro Leu
 65 70 75 80

 Ser Val Leu Ile Leu Gly Leu Leu Ile Met Phe Ile Thr Ile Val Phe
 85 90 95

 Cys Ala Ala Gly Ile Val Val Leu Val His Arg Arg Arg Arg Tyr
 100 105 110

 Lys Lys Lys Gln Val Asp Glu Gln Met Arg Asp Asn Ser Pro Val His
 115 120 125

Leu Gln Tyr Ser Met Tyr Gly His Lys Thr Thr His His Thr Thr Glu
 130 135 140
 Arg Pro Ser Ala Ser Leu Tyr Glu Gln His Met Val Ser Pro Met Val
 145 150 155 160
 His Val Tyr Arg Ser Pro Ser Phe Gly Pro Lys His Leu Glu Glu Glu
 165 170 175
 Glu Glu Arg Asn Glu Lys Glu Gly Ser Asp Ala Lys His Leu Gln Arg
 180 185 190
 Ser Leu Leu Glu Gln Glu Asn His Ser Pro Leu Thr Gly Ser Asn Met
 195 200 205
 Lys Tyr Lys Thr Thr Asn Gln Ser Thr Glu Phe Leu Ser Phe Gln Asp
 210 215 220
 Ala Ser Ser Leu Tyr Arg Asn Ile Leu Glu Lys Glu Arg Glu Leu Gln
 225 230 235 240
 Gln Leu Gly Ile Thr Glu Tyr Leu Arg Lys Asn Ile Ala Gln Leu Gln
 245 250 255
 Pro Asp Met Glu Ala His Tyr Pro Gly Ala His Glu Glu Leu Lys Leu
 260 265 270
 Met Glu Thr Leu Met Tyr Ser Arg Pro Arg Lys Val Leu Val Glu Gln
 275 280 285
 Thr Lys Asn Glu Tyr Phe Glu Leu Lys Ala Asn Leu His Ala Glu Pro
 290 295 300
 Asp Tyr Leu Glu Val Leu Glu Gln Gln Thr
 305 310

<210> 32
 <211> 966
 <212> PRT
 <213> Homo sapiens

<400> 32
 Arg Arg Gly Ala Gln Gly Gly Lys Met His Thr Cys Cys Pro Pro Val
 1 5 10 15
 Thr Leu Glu Gln Asp Leu His Arg Lys Met His Ser Trp Met Leu Gln
 20 25 30
 Thr Leu Ala Phe Ala Val Thr Ser Leu Val Leu Ser Cys Ala Glu Thr
 35 40 45
 Ile Asp Tyr Tyr Gly Glu Ile Cys Asp Asn Ala Cys Pro Cys Glu Glu
 50 55 60
 Lys Asp Gly Ile Leu Thr Val Ser Cys Glu Asn Arg Gly Ile Ile Ser

65

70

75

80

Leu Ser Glu Ile Ser Pro Pro Arg Phe Pro Ile Tyr His Leu Leu			
85	90	95	
Ser Gly Asn Leu Leu Asn Arg Leu Tyr Pro Asn Glu Phe Val Asn Tyr			
100	105	110	
Thr Gly Ala Ser Ile Leu His Leu Gly Ser Asn Val Ile Gln Asp Ile			
115	120	125	
Glu Thr Gly Ala Phe His Gly Leu Arg Gly Leu Arg Arg Leu His Leu			
130	135	140	
Asn Asn Asn Lys Leu Glu Leu Leu Arg Asp Asp Thr Phe Leu Gly Leu			
145	150	155	160
Glu Asn Leu Glu Tyr Leu Gln Val Asp Tyr Asn Tyr Ile Ser Val Ile			
165	170	175	
Glu Pro Asn Ala Phe Gly Lys Leu His Leu Leu Gln Val Leu Ile Leu			
180	185	190	
Asn Asp Asn Leu Leu Ser Ser Leu Pro Asn Asn Leu Phe Arg Phe Val			
195	200	205	
Pro Leu Thr His Leu Asp Leu Arg Gly Asn Arg Leu Lys Leu Leu Pro			
210	215	220	
Tyr Val Gly Leu Leu Gln His Met Asp Lys Val Val Glu Leu Gln Leu			
225	230	235	240
Glu Glu Asn Pro Trp Asn Cys Ser Cys Glu Leu Ile Ser Leu Lys Asp			
245	250	255	
Trp Leu Asp Ser Ile Ser Tyr Ser Ala Leu Val Gly Asp Val Val Cys			
260	265	270	
Glu Thr Pro Phe Arg Leu His Gly Arg Asp Leu Asp Glu Val Ser Lys			
275	280	285	
Gln Glu Leu Cys Pro Arg Arg Leu Ile Ser Asp Tyr Glu Met Arg Pro			
290	295	300	
Gln Thr Pro Leu Ser Thr Thr Gly Tyr Leu His Thr Thr Pro Ala Ser			
305	310	315	320
Val Asn Ser Val Ala Thr Ser Ser Ala Val Tyr Lys Pro Pro Leu			
325	330	335	
Lys Pro Pro Lys Gly Thr Arg Gln Pro Asn Lys Pro Arg Val Arg Pro			
340	345	350	
Thr Ser Arg Gln Pro Ser Lys Asp Leu Gly Tyr Ser Asn Tyr Gly Pro			
355	360	365	
Ser Ile Ala Tyr Gln Thr Lys Ser Pro Val Pro Leu Glu Cys Pro Thr			

370

375

380

Ala Cys Ser Cys Asn Leu Gln Ile Ser Asp Leu Gly Leu Asn Val Asn
 385 390 395 400

Cys Gln Glu Arg Lys Ile Glu Ser Ile Ala Glu Leu Gln Pro Lys Pro
 405 410 415

Tyr Asn Pro Lys Lys Met Tyr Leu Thr Glu Asn Tyr Ile Ala Val Val
 420 425 430

Arg Arg Thr Asp Phe Leu Glu Ala Thr Gly Leu Asp Leu Leu His Leu
 435 440 445

Gly Asn Asn Arg Ile Ser Met Ile Gln Asp Arg Ala Phe Gly Asp Leu
 450 455 460

Thr Asn Leu Arg Arg Leu Tyr Leu Asn Gly Asn Arg Ile Glu Arg Leu
 465 470 475 480

Ser Pro Glu Leu Phe Tyr Gly Leu Gln Ser Leu Gln Tyr Leu Phe Leu
 485 490 495

Gln Tyr Asn Leu Ile Arg Glu Ile Gln Ser Gly Thr Phe Asp Pro Val
 500 505 510

Pro Asn Leu Gln Leu Leu Phe Leu Asn Asn Asn Leu Leu Gln Ala Met
 515 520 525

Pro Ser Gly Val Phe Ser Gly Leu Thr Leu Leu Arg Leu Asn Leu Arg
 530 535 540

Ser Asn His Phe Thr Ser Leu Pro Val Ser Gly Val Leu Asp Gln Leu
 545 550 555 560

Lys Ser Leu Ile Gln Ile Asp Leu His Asp Asn Pro Trp Asp Cys Thr
 565 570 575

Cys Asp Ile Val Gly Met Lys Leu Trp Val Glu Gln Leu Lys Val Gly
 580 585 590

Val Leu Val Asp Glu Val Ile Cys Lys Ala Pro Lys Lys Phe Ala Glu
 595 600 605

Thr Asp Met Arg Ser Ile Lys Ser Glu Leu Leu Cys Pro Asp Tyr Ser
 610 615 620

Asp Val Val Val Ser Thr Pro Thr Pro Ser Ser Ile Gln Val Pro Ala
 625 630 635 640

Arg Thr Ser Ala Val Thr Pro Ala Val Arg Leu Asn Ser Thr Gly Ala
 645 650 655

Pro Ala Ser Leu Gly Ala Gly Gly Ala Ser Ser Val Pro Leu Ser
 660 665 670

Val Leu Ile Leu Ser Leu Leu Val Phe Ile Met Ser Val Phe Val

675

680

685

Ala Ala Gly Leu Phe Val Leu Val Met Lys Arg Arg Lys Lys Asn Gln
 690 695 700

Ser Asp His Thr Ser Thr Asn Asn Ser Asp Val Ser Ser Phe Asn Met
 705 710 715 720

Gln Tyr Ser Val Tyr Gly Gly Gly Gly Thr Gly Gly His Pro His
 725 730 735

Ala His Val His His Arg Gly Pro Ala Leu Pro Lys Val Lys Thr Pro
 740 745 750

Ala Gly His Val Tyr Glu Tyr Ile Pro His Pro Leu Gly His Met Cys
 755 760 765

Lys Asn Pro Ile Tyr Arg Ser Arg Glu Gly Asn Ser Val Glu Asp Tyr
 770 775 780

Lys Asp Leu His Glu Leu Lys Val Thr Tyr Ser Ser Asn His His Leu
 785 790 795 800

Gln Gln Gln Gln Pro Pro Pro Pro Gln Gln Pro Gln Gln Gln
 805 810 815

Pro Pro Pro Gln Leu Gln Leu Gln Pro Gly Glu Glu Glu Arg Arg Glu
 820 825 830

Ser His His Leu Arg Ser Pro Ala Tyr Ser Val Ser Thr Ile Glu Pro
 835 840 845

Arg Glu Asp Leu Leu Ser Pro Val Gln Asp Ala Asp Arg Phe Tyr Arg
 850 855 860

Gly Ile Leu Glu Pro Asp Lys His Cys Ser Thr Thr Pro Ala Gly Asn
 865 870 875 880

Ser Leu Pro Glu Tyr Pro Lys Phe Pro Cys Ser Pro Ala Ala Tyr Thr
 885 890 895

Phe Ser Pro Asn Tyr Asp Leu Arg Arg Pro His Gln Tyr Leu His Pro
 900 905 910

Gly Ala Gly Asp Ser Arg Leu Arg Glu Pro Val Leu Tyr Ser Pro Pro
 915 920 925

Ser Ala Val Phe Val Glu Pro Asn Arg Asn Glu Tyr Leu Glu Leu Lys
 930 935 940

Ala Lys Leu Asn Val Glu Pro Asp Tyr Leu Glu Val Leu Glu Lys Gln
 945 950 955 960

Thr Thr Phe Ser Gln Phe
 965

<210> 33
<211> 1428
<212> PRT
<213> Mus musculus

<400> 33
Met Val Ala Val Ala Val Ala Ala Ser Thr Glu Ala Arg Leu Arg Gly
1 5 10 15

Ser Thr Thr Ala Thr Ala Ala Pro Ala Gly Arg Lys Gly Arg Gln His
20 25 30

Arg Pro Cys Thr Ala Thr Gly Ala Trp Arg Pro Gly Pro Arg Ala Arg
35 40 45

Leu Cys Leu Pro Arg Val Leu Ser Arg Ala Leu Pro Pro Pro Pro Leu
50 55 60

Leu Pro Leu Leu Phe Ser Leu Leu Leu Pro Leu Pro Arg Glu Ala
65 70 75 80

Glu Ala Ala Ala Val Ala Ala Ala Val Ser Gly Ser Ala Ala Ala Glu
85 90 95

Ala Lys Glu Cys Asp Arg Pro Cys Val Asn Gly Gly Arg Cys Asn Pro
100 105 110

Gly Thr Gly Gln Cys Val Cys Pro Thr Gly Trp Val Gly Glu Gln Cys
115 120 125

Gln His Cys Gly Gly Arg Phe Arg Leu Thr Gly Ser Ser Gly Phe Val
130 135 140

Thr Asp Gly Pro Gly Asn Tyr Lys Tyr Lys Thr Lys Cys Thr Trp Leu
145 150 155 160

Ile Glu Gly Tyr Pro Asn Ala Val Leu Arg Leu Arg Phe Asn His Phe
165 170 175

Ala Thr Glu Cys Ser Trp Asp His Leu Tyr Val Tyr Asp Gly Asp Ser
180 185 190

Ile Tyr Ala Pro Leu Ile Ala Ala Phe Ser Gly Leu Ile Val Pro Glu
195 200 205

Arg Asp Gly Asn Glu Thr Ala Pro Glu Val Thr Val Thr Ser Gly Tyr
210 215 220

Ala Leu Leu His Phe Phe Ser Asp Ala Ala Tyr Asn Leu Thr Gly Phe
225 230 235 240

Asn Ile Thr Tyr Asn Phe Asp Met Cys Pro Asn Asn Cys Ser Ala Arg
245 250 255

Gly Glu Cys Lys Ser Ser Asn Ser Ser Ala Val Glu Cys Glu Cys
260 265 270

Ser Glu Asn Trp Lys Gly Glu Ser Cys Asp Ile Pro His Cys Thr Asp
 275 280 285

 Asn Cys Gly Phe Pro His Arg Gly Ile Cys Asn Ala Ser Asp Thr Arg
 290 295 300

 Gly Cys Ser Cys Phe Pro His Trp Gln Gly Pro Gly Cys Ser Ile Pro
 305 310 315 320

 Val Pro Ala Asn Gln Ser Phe Trp Thr Arg Glu Glu Tyr Ser Asp Leu
 325 330 335

 Lys Leu Pro Arg Ala Ser His Lys Ala Val Val Asn Gly Asn Ile Met
 340 345 350

 Trp Val Val Gly Gly Tyr Met Phe Asn His Ser Asp Tyr Ser Met Val
 355 360 365

 Leu Ala Tyr Asp Leu Thr Ser Arg Glu Trp Leu Pro Leu Asn His Ser
 370 375 380

 Val Asn Ser Val Val Val Arg Tyr Gly His Ser Leu Ala Leu His Lys
 385 390 395 400

 Asp Lys Ile Tyr Met Tyr Gly Gly Lys Ile Asp Ser Thr Gly Asn Val
 405 410 415

 Thr Asn Glu Leu Arg Val Phe His Ile His Asn Glu Ser Trp Val Leu
 420 425 430

 Leu Thr Pro Lys Ala Lys Asp Gln Tyr Ala Val Val Gly His Ser Ala
 435 440 445

 His Ile Val Thr Leu Ala Ser Gly Arg Val Val Met Leu Val Ile Phe
 450 455 460

 Gly His Cys Pro Leu Tyr Gly Tyr Ile Ser Val Val Gln Glu Tyr Asp
 465 470 475 480

 Leu Glu Lys Asn Thr Trp Ser Ile Leu His Thr Gln Gly Ala Leu Val
 485 490 495

 Gln Gly Gly Tyr Gly His Ser Ser Ala Tyr Asp Asp Arg Thr Lys Ala
 500 505 510

 Leu Tyr Val His Gly Gly Tyr Lys Ala Phe Ser Ala Asn Lys Tyr Arg
 515 520 525

 Leu Ala Asp Asp Leu Tyr Arg Tyr Asp Val Asp Thr Gln Met Trp Thr
 530 535 540

 Ile Leu Lys Asp Ser Arg Phe Phe Arg Tyr Leu His Thr Ala Val Ile
 545 550 555 560

 Val Ser Gly Thr Met Leu Val Phe Gly Gly Asn Thr His Asn Asp Thr
 565 570 575

Ser Met Ser His Gly Ala Lys Cys Phe Ser Ser Asp Phe Met Ala Tyr
 580 585 590
 Asp Ile Ala Cys Asp Arg Trp Ser Val Leu Pro Arg Pro Glu Leu His
 595 600 605
 His Asp Val Asn Arg Phe Gly His Ser Ala Val Leu Tyr Asn Ser Thr
 610 615 620
 Met Tyr Val Phe Gly Gly Phe Asn Ser Leu Leu Leu Ser Asp Val Leu
 625 630 635 640
 Val Phe Thr Ser Glu Gln Cys Asp Ala His Arg Ser Glu Ala Ala Cys
 645 650 655
 Val Ala Ala Gly Pro Gly Ile Arg Cys Leu Trp Asp Thr Gln Ser Ser
 660 665 670
 Arg Cys Thr Ser Trp Glu Leu Ala Thr Glu Glu Gln Ala Glu Lys Leu
 675 680 685
 Lys Ser Glu Cys Phe Ser Lys Arg Thr Leu Asp His Asp Arg Cys Asp
 690 695 700
 Gln His Thr Asp Cys Tyr Ser Cys Thr Ala Asn Thr Asn Asp Cys His
 705 710 715 720
 Trp Cys Asn Asp His Cys Val Pro Val Asn His Ser Cys Thr Glu Gly
 725 730 735
 Gln Ile Ser Ile Ala Lys Tyr Glu Ser Cys Pro Lys Asp Asn Pro Met
 740 745 750
 Tyr Tyr Cys Asn Lys Lys Thr Ser Cys Arg Ser Cys Ala Leu Asp Gln
 755 760 765
 Asn Cys Gln Trp Glu Pro Arg Asn Gln Glu Cys Ile Ala Leu Pro Glu
 770 775 780
 Asn Ile Cys Gly Asn Gly Trp His Leu Val Gly Asn Ser Cys Leu Lys
 785 790 795 800
 Ile Thr Thr Ala Lys Glu Asn Tyr Asp Asn Ala Lys Leu Ser Cys Arg
 805 810 815
 Asn His Asn Ala Phe Leu Ala Ser Leu Thr Ser Gln Lys Lys Val Glu
 820 825 830
 Leu Val Leu Lys Gln Leu Arg Leu Met Gln Ser Ser Gln Ser Met Ser
 835 840 845
 Lys Leu Thr Leu Thr Pro Trp Val Gly Leu Arg Lys Ile Asn Val Ser
 850 855 860
 Tyr Trp Cys Trp Glu Asp Met Ser Pro Phe Thr Asn Ser Leu Leu Gln
 865 870 875 880

Trp Met Pro Ser Glu Pro Ser Asp Ala Gly Phe Cys Gly Ile Leu Ser
 885 890 895

 Glu Pro Ser Thr Arg Gly Leu Lys Ala Ala Thr Cys Ile Asn Pro Leu
 900 905 910

 Asn Gly Ser Val Cys Glu Arg Pro Ala Asn His Ser Ala Lys Gln Cys
 915 920 925

 Arg Thr Pro Cys Ala Leu Arg Thr Ala Cys Gly Glu Cys Thr Ser Ser
 930 935 940

 Ser Ser Glu Cys Met Trp Cys Ser Asn Met Lys Gln Cys Val Asp Ser
 945 950 955 960

 Asn Ala Tyr Val Ala Ser Phe Pro Phe Gly Gln Cys Met Glu Trp Tyr
 965 970 975

 Thr Met Ser Ser Cys Pro Pro Glu Asn Cys Ser Gly Tyr Cys Thr Cys
 980 985 990

 Ser His Cys Leu Glu Gln Pro Gly Cys Gly Trp Cys Thr Asp Pro Ser
 995 1000 1005

 Asn Thr Gly Lys Gly Lys Cys Ile Glu Gly Ser Tyr Lys Gly Pro Val
 1010 1015 1020

 Lys Met Pro Ser Gln Ala Ser Ala Gly Asn Val Tyr Pro Gln Pro Leu
 1025 1030 1035 1040

 Leu Asn Ser Ser Met Cys Leu Glu Asp Ser Arg Tyr Asn Trp Ser Phe
 1045 1050 1055

 Ile His Cys Pro Ala Cys Gln Cys Asn Gly His Ser Lys Cys Ile Asn
 1060 1065 1070

 Gln Ser Val Cys Glu Lys Cys Glu Asp Leu Thr Thr Gly Lys His Cys
 1075 1080 1085

 Glu Thr Cys Ile Ser Gly Phe Tyr Gly Asp Pro Thr Asn Gly Gly Lys
 1090 1095 1100

 Cys Gln Pro Cys Lys Cys Asn Gly His Ala Ser Leu Cys Asn Thr Asn
 1105 1110 1115 1120

 Thr Gly Lys Cys Phe Cys Thr Thr Lys Gly Val Lys Gly Asp Glu Cys
 1125 1130 1135

 Gln Leu Cys Lys Val Glu Asn Arg Tyr Gln Gly Asn Pro Leu Lys Gly
 1140 1145 1150

 Thr Cys Tyr Tyr Thr Leu Leu Ile Asp Tyr Gln Phe Thr Phe Ser Leu
 1155 1160 1165

 Ser Gln Gly Asp Asp Arg Tyr Tyr Thr Ala Ile Asn Phe Val Ala Thr
 1170 1175 1180

Pro Asp Glu Gln Asn Arg Asp Phe Asp Met Phe Ile Asn Ala Ser Lys
 1185 1190 1195 1200

 Lys Phe Asn Leu Asn Ile Thr Trp Ala Thr Ser Phe Pro Ala Gly Thr
 1205 1210 1215

 Gln Thr Gly Glu Glu Val Pro Val Val Ser Lys Thr Asn Ile Lys Glu
 1220 1225 1230

 Tyr Lys Asp Ser Phe Ser Asn Glu Lys Phe Asp Phe Arg Asn His Pro
 1235 1240 1245

 Asn Ile Thr Phe Phe Val Tyr Val Ser Asn Phe Thr Trp Pro Ile Lys
 1250 1255 1260

 Ile Gln Ile Ala Phe Ser Gln His Ser Asn Phe Met Asp Leu Val Gln
 1265 1270 1275 1280

 Phe Phe Val Thr Phe Phe Ser Cys Phe Leu Ser Leu Leu Val Ala
 1285 1290 1295

 Ala Val Val Trp Lys Ile Lys Gln Ser Cys Trp Ala Ser Arg Arg Arg
 1300 1305 1310

 Glu Gln Leu Leu Arg Glu Met Gln Gln Met Ala Ser Arg Pro Phe Ala
 1315 1320 1325

 Ser Val Asn Val Ala Leu Glu Thr Asp Glu Glu Pro Pro Asp Leu Ile
 1330 1335 1340

 Gly Gly Ser Ile Lys Thr Val Pro Lys Pro Ile Ala Leu Glu Pro Cys
 1345 1350 1355 1360

 Phe Gly Asn Lys Ala Ala Val Leu Ser Val Phe Val Arg Leu Pro Arg
 1365 1370 1375

 Gly Leu Gly Gly Ile Pro Pro Pro Gly Gln Ser Gly Leu Ala Val Ala
 1380 1385 1390

 Ser Ala Leu Val Asp Ile Ser Gln Gln Met Pro Ile Val Tyr Lys Glu
 1395 1400 1405

 Lys Ser Gly Ala Val Arg Asn Arg Lys Gln Gln Pro Pro Ala Gln Pro
 1410 1415 1420

 Gly Thr Cys Ile
 1425

 <210> 34
 <211> 1428
 <212> PRT
 <213> Mus musculus

 <400> 34
 Met Val Ala Val Ala Ala Ala Ala Thr Glu Ala Arg Leu Arg Gly
 1 5 10 15

Ser Thr Thr Thr Ala Ala Pro Ala Gly Arg Lys Gly Arg Gln His
 20 25 30

 Arg Pro Cys Thr Ala Thr Gly Ala Trp Arg Pro Gly Pro Arg Ala Arg
 35 40 45

 Leu Cys Leu Pro Arg Val Leu Ser Arg Ala Leu Pro Pro Pro Pro Leu
 50 55 60

 Leu Pro Leu Leu Phe Ser Leu Leu Leu Pro Leu Pro Arg Glu Ala
 65 70 75 80

 Glu Ala Ala Ala Val Ala Ala Ala Val Ser Gly Ser Ala Ala Ala Glu
 85 90 95

 Ala Lys Glu Cys Asp Arg Pro Cys Val Asn Gly Gly Arg Cys Asn Pro
 100 105 110

 Gly Thr Gly Gln Cys Val Cys Pro Thr Gly Trp Val Gly Glu Gln Cys
 115 120 125

 Gln His Cys Gly Gly Arg Phe Arg Leu Thr Gly Ser Ser Gly Phe Val
 130 135 140

 Thr Asp Gly Pro Gly Asn Tyr Lys Tyr Lys Thr Lys Cys Thr Trp Leu
 145 150 155 160

 Ile Glu Gly Gln Pro Asn Arg Ile Met Arg Leu Arg Phe Asn His Phe
 165 170 175

 Ala Thr Glu Cys Ser Trp Asp His Leu Tyr Val Tyr Asp Gly Asp Ser
 180 185 190

 Ile Tyr Ala Pro Leu Ile Ala Ala Phe Ser Gly Leu Ile Val Pro Glu
 195 200 205

 Arg Asp Gly Asn Glu Thr Ala Pro Glu Val Thr Val Thr Ser Gly Tyr
 210 215 220

 Ala Leu Leu His Phe Phe Ser Asp Ala Ala Tyr Asn Leu Thr Gly Phe
 225 230 235 240

 Asn Ile Thr Tyr Asn Phe Asp Met Cys Pro Asn Asn Cys Ser Gly Arg
 245 250 255

 Gly Glu Cys Lys Ser Ser Asn Ser Ser Ser Ala Val Glu Cys Glu Cys
 260 265 270

 Ser Glu Asn Trp Lys Gly Glu Ser Cys Asp Ile Pro His Cys Thr Asp
 275 280 285

 Asn Cys Gly Phe Pro His Arg Gly Ile Cys Asn Ala Ser Asp Thr Arg
 290 295 300

 Gly Cys Ser Cys Phe Pro His Trp Gln Gly Pro Gly Cys Ser Ile Pro
 305 310 315 320

Val Pro Ala Asn Gln Ser Phe Trp Thr Arg Glu Glu Tyr Ser Asp Leu
 325 330 335

 Lys Leu Pro Arg Ala Ser His Lys Ala Val Val Asn Gly Asn Ile Met
 340 345 350

 Trp Val Val Gly Gly Tyr Met Phe Asn His Ser Asp Tyr Ser Met Val
 355 360 365

 Leu Ala Tyr Asp Leu Thr Ser Arg Glu Trp Leu Pro Leu Asn His Ser
 370 375 380

 Val Asn Ser Val Val Val Arg Tyr Gly His Ser Leu Ala Leu His Lys
 385 390 395 400

 Asp Lys Ile Tyr Met Tyr Gly Gly Lys Ile Asp Ser Thr Gly Asn Val
 405 410 415

 Thr Asn Glu Leu Arg Val Phe His Ile His Asn Glu Ser Trp Val Leu
 420 425 430

 Leu Thr Pro Lys Ala Lys Asp Gln Tyr Ala Val Val Gly His Ser Ala
 435 440 445

 His Ile Val Thr Leu Ala Ser Gly Arg Val Val Met Leu Val Ile Phe
 450 455 460

 Gly His Cys Pro Leu Tyr Gly Tyr Ile Ser Val Val Gln Glu Tyr Asp
 465 470 475 480

 Leu Glu Lys Asn Thr Trp Ser Ile Leu His Thr Gln Gly Ala Leu Val
 485 490 495

 Gln Gly Gly Tyr Gly His Ser Ser Val Tyr Asp Asp Arg Thr Lys Ala
 500 505 510

 Leu Tyr Val His Gly Gly Tyr Lys Ala Phe Ser Ala Asn Lys Tyr Arg
 515 520 525

 Leu Ala Asp Asp Leu Tyr Arg Tyr Asp Val Asp Thr Gln Met Trp Thr
 530 535 540

 Ile Leu Lys Asp Ser Arg Phe Phe Arg Tyr Leu His Thr Ala Val Ile
 545 550 555 560

 Val Ser Gly Thr Met Leu Val Phe Gly Gly Asn Thr His Asn Asp Thr
 565 570 575

 Ser Met Ser His Gly Ala Lys Cys Phe Ser Ser Asp Phe Met Ala Tyr
 580 585 590

 Asp Ile Ala Cys Asp Arg Trp Ser Val Leu Pro Arg Pro Glu Leu His
 595 600 605

 His Asp Val Asn Arg Phe Gly His Ser Ala Val Leu Tyr Asn Ser Thr
 610 615 620

Met Tyr Val Phe Gly Gly Phe Asn Ser Leu Leu Leu Ser Asp Val Leu
 625 630 635 640

Val Phe Thr Ser Glu Gln Cys Asp Ala His Arg Ser Glu Ala Ala Cys
 645 650 655

Val Ala Ala Gly Pro Gly Ile Arg Cys Leu Trp Asp Thr Gln Ser Ser
 660 665 670

Arg Cys Thr Ser Trp Glu Leu Ala Thr Glu Glu Gln Ala Glu Lys Leu
 675 680 685

Lys Ser Glu Cys Phe Ser Lys Arg Thr Leu Asp His Asp Arg Cys Asp
 690 695 700

Gln His Thr Asp Cys Tyr Ser Cys Thr Ala Asn Thr Asn Asp Cys His
 705 710 715 720

Trp Cys Asn Asp His Cys Val Pro Val Asn His Ser Cys Thr Glu Gly
 725 730 735

Gln Ile Ser Ile Ala Lys Tyr Glu Ser Cys Pro Lys Asp Asn Pro Met
 740 745 750

Tyr Tyr Cys Asn Lys Lys Thr Ser Cys Arg Ser Cys Ala Leu Asp Gln
 755 760 765

Asn Cys Gln Trp Glu Pro Arg Asn Gln Glu Cys Ile Ala Leu Pro Glu
 770 775 780

Asn Ile Cys Gly Asn Gly Trp His Leu Val Gly Asn Ser Cys Leu Lys
 785 790 795 800

Ile Thr Thr Ala Lys Glu Asn Tyr Asp Asn Ala Lys Leu Ser Cys Arg
 805 810 815

Asn His Asn Ala Phe Leu Ala Ser Leu Thr Ser Gln Lys Lys Val Glu
 820 825 830

Phe Val Leu Lys Gln Leu Arg Leu Met Gln Ser Ser Gln Ser Met Ser
 835 840 845

Lys Leu Thr Leu Thr Pro Trp Val Gly Leu Arg Lys Ile Asn Val Ser
 850 855 860

Tyr Trp Cys Trp Glu Asp Met Ser Pro Phe Thr Asn Ser Leu Leu Gln
 865 870 875 880

Trp Met Pro Ser Glu Pro Ser Asp Ala Gly Phe Cys Gly Ile Leu Ser
 885 890 895

Glu Pro Ser Thr Arg Gly Leu Lys Ala Ala Thr Cys Ile Asn Pro Leu
 900 905 910

Asn Gly Ser Val Cys Glu Arg Pro Ala Asn His Ser Ala Lys Gln Cys
 915 920 925

Arg Thr Pro Cys Ala Leu Arg Thr Ala Cys Gly Glu Cys Thr Ser Ser
 930 935 940

Ser Ser Glu Cys Met Trp Cys Ser Asn Met Lys Gln Cys Val Asp Ser
 945 950 955 960

Asn Ala Tyr Val Ala Ser Phe Pro Phe Gly Gln Cys Met Glu Trp Tyr
 965 970 975

Thr Met Ser Ser Cys Pro Pro Glu Asn Cys Ser Gly Tyr Cys Thr Cys
 980 985 990

Ser His Cys Leu Glu Gln Pro Gly Cys Gly Trp Cys Thr Asp Pro Ser
 995 1000 1005

Asn Thr Gly Lys Gly Lys Cys Ile Glu Gly Ser Tyr Lys Gly Pro Val
 1010 1015 1020

Lys Met Pro Ser Gln Ala Ser Ala Gly Asn Val Tyr Pro Gln Pro Leu
 1025 1030 1035 1040

Leu Asn Ser Ser Met Cys Leu Glu Asp Ser Arg Tyr Asn Trp Ser Phe
 1045 1050 1055

Ile His Cys Pro Ala Cys Gln Cys Asn Gly His Ser Lys Cys Ile Asn
 1060 1065 1070

Gln Ser Ile Cys Glu Lys Cys Glu Asp Leu Thr Thr Gly Lys His Cys
 1075 1080 1085

Glu Thr Cys Ile Ser Gly Phe Tyr Gly Asp Pro Thr Asn Gly Gly Lys
 1090 1095 1100

Cys Gln Pro Cys Lys Cys Asn Gly His Ala Ser Leu Cys Asn Thr Asn
 1105 1110 1115 1120

Thr Gly Lys Cys Phe Cys Thr Thr Lys Gly Val Lys Gly Asp Glu Cys
 1125 1130 1135

Gln Leu Cys Glu Val Glu Asn Arg Tyr Gln Gly Asn Pro Leu Lys Gly
 1140 1145 1150

Thr Cys Tyr Tyr Thr Leu Leu Ile Asp Tyr Gln Phe Thr Phe Ser Leu
 1155 1160 1165

Ser Gln Glu Asp Asp Arg Tyr Tyr Thr Ala Ile Asn Phe Val Ala Thr
 1170 1175 1180

Pro Asp Glu Gln Asn Arg Asp Leu Asp Met Phe Ile Asn Ala Ser Lys
 1185 1190 1195 1200

Asn Phe Asn Leu Asn Ile Thr Trp Ala Thr Ser Phe Pro Ala Gly Thr
 1205 1210 1215

Gln Thr Gly Glu Glu Val Pro Val Val Ser Lys Thr Asn Ile Lys Glu
 1220 1225 1230

Tyr Lys Asp Ser Phe Ser Asn Glu Lys Phe Asp Phe Arg Asn His Pro
 1235 1240 1245
 Asn Ile Thr Phe Phe Val Tyr Val Ser Asn Phe Thr Trp Pro Ile Lys
 1250 1255 1260
 Ile Gln Ile Ala Phe Ser Gln His Ser Asn Phe Met Asp Leu Val Gln
 1265 1270 1275 1280
 Phe Phe Val Thr Phe Phe Ser Cys Phe Leu Ser Leu Leu Val Ala
 1285 1290 1295
 Ala Val Val Trp Lys Ile Lys Gln Ser Cys Trp Ala Ser Arg Arg Arg
 1300 1305 1310
 Glu Gln Leu Leu Arg Glu Met Gln Gln Met Ala Ser Arg Pro Phe Ala
 1315 1320 1325
 Ser Val Asn Val Ala Leu Glu Thr Asp Glu Glu Pro Pro Asp Leu Ile
 1330 1335 1340
 Gly Gly Ser Ile Lys Thr Val Pro Lys Pro Ile Ala Leu Glu Pro Cys
 1345 1350 1355 1360
 Phe Gly Asn Lys Ala Ala Val Leu Ser Val Phe Val Arg Leu Pro Arg
 1365 1370 1375
 Gly Leu Gly Gly Ile Pro Pro Pro Gly Gln Ser Gly Leu Ala Val Ala
 1380 1385 1390
 Ser Ala Leu Val Asp Ile Ser Gln Gln Met Pro Ile Val Tyr Lys Glu
 1395 1400 1405
 Lys Ser Gly Ala Val Arg Asn Arg Lys Gln Gln Pro Pro Ala Gln Pro
 1410 1415 1420
 Gly Thr Cys Ile
 1425

<210> 35
 <211> 1198
 <212> PRT
 <213> Homo sapiens

<400> 35
 Met Val Ala Ala Ala Ala Ala Thr Glu Ala Arg Leu Arg Arg Arg Thr
 1 5 10 15
 Ala Ala Thr Ala Ala Leu Ala Gly Arg Ser Gly Gly Pro His Cys Val
 20 25 30
 Asn Gly Gly Arg Cys Asn Pro Gly Thr Gly Gln Cys Val Cys Pro Ala
 35 40 45
 Gly Trp Val Gly Glu Gln Cys Gln His Cys Gly Gly Arg Phe Arg Leu

50	55	60
Thr Gly Ser Ser Gly Phe Val Thr Asp Gly Pro Gly Asn Tyr Lys Tyr		
65	70	75
80		
Lys Thr Lys Cys Thr Trp Leu Ile Glu Gly Gln Pro Asn Arg Ile Met		
85	90	95
Arg Leu Arg Phe Asn His Phe Ala Thr Glu Cys Ser Trp Asp His Leu		
100	105	110
Tyr Val Tyr Asp Gly Asp Ser Ile Tyr Ala Pro Leu Val Ala Ala Phe		
115	120	125
Ser Gly Leu Ile Val Pro Glu Arg Asp Gly Asn Glu Thr Val Pro Glu		
130	135	140
Val Val Ala Thr Ser Gly Tyr Ala Leu Leu His Phe Phe Ser Asp Ala		
145	150	155
160		
Ala Tyr Asn Leu Thr Gly Phe Asn Ile Thr Tyr Ser Phe Asp Met Cys		
165	170	175
Pro Asn Asn Cys Ser Gly Arg Gly Glu Cys Lys Ile Ser Asn Ser Ser		
180	185	190
Asp Thr Val Glu Cys Glu Cys Ser Glu Asn Trp Lys Gly Glu Ala Cys		
195	200	205
Asp Ile Pro His Cys Thr Asp Asn Cys Gly Phe Pro His Arg Gly Ile		
210	215	220
Cys Asn Ser Ser Asp Val Arg Gly Cys Ser Cys Phe Ser Asp Trp Gln		
225	230	235
240		
Gly Pro Gly Cys Ser Val Pro Val Pro Ala Asn Gln Ser Phe Trp Thr		
245	250	255
Arg Glu Glu Tyr Ser Asn Leu Lys Leu Pro Arg Ala Ser His Lys Ala		
260	265	270
Val Val Asn Gly Asn Ile Met Trp Val Val Gly Gly Tyr Met Phe Asn		
275	280	285
His Ser Asp Tyr Asn Met Val Leu Ala Tyr Asp Leu Ala Ser Arg Glu		
290	295	300
Trp Leu Pro Leu Asn Arg Ser Val Asn Asn Val Val Val Arg Tyr Gly		
305	310	315
320		
His Ser Leu Ala Leu Tyr Lys Asp Lys Ile Tyr Met Tyr Gly Gly Lys		
325	330	335
Ile Asp Ser Thr Gly Asn Val Thr Asn Glu Leu Arg Val Phe His Ile		
340	345	350
His Asn Glu Ser Trp Val Leu Leu Thr Pro Lys Ala Lys Glu Gln Tyr		

355	360	365
Ala Val Val Gly His Ser Ala His Ile Val Thr Leu Lys Asn Gly Arg		
370	375	380
Val Val Met Leu Val Ile Phe Gly His Cys Pro Leu Tyr Gly Tyr Ile		
385	390	395
Ser Asn Val Gln Glu Tyr Asp Leu Asp Lys Asn Thr Trp Ser Ile Leu		
405	410	415
His Thr Gln Gly Ala Leu Val Gln Gly Gly Tyr Gly His Ser Ser Val		
420	425	430
Tyr Asp His Arg Thr Arg Ala Leu Tyr Val His Gly Gly Tyr Lys Ala		
435	440	445
Phe Ser Ala Asn Lys Tyr Arg Leu Ala Asp Asp Leu Tyr Arg Tyr Asp		
450	455	460
Val Asp Thr Gln Met Trp Thr Ile Leu Lys Asp Ser Arg Phe Phe Arg		
465	470	475
Tyr Leu His Thr Ala Val Ile Val Ser Gly Thr Met Leu Val Phe Gly		
485	490	495
Gly Asn Thr His Asn Asp Thr Ser Met Ser His Gly Ala Lys Cys Phe		
500	505	510
Ser Ser Asp Phe Met Ala Tyr Asp Ile Ala Cys Asp Arg Trp Ser Val		
515	520	525
Leu Pro Arg Pro Asp Ser Thr Met Met Ser Thr Asp Leu Ala Ile Pro		
530	535	540
Ala Val Leu His Asn Ser Thr Met Tyr Val Phe Gly Gly Phe Asn Ser		
545	550	555
Leu Leu Leu Ser Asp Ile Leu Val Phe Thr Ser Glu Gln Cys Asp Ala		
565	570	575
His Arg Ser Glu Ala Ala Cys Leu Ala Ala Gly Pro Gly Ile Arg Cys		
580	585	590
Val Trp Asn Thr Gly Ser Ser Gln Cys Ile Ser Trp Ala Leu Ala Thr		
595	600	605
Asp Glu Gln Glu Glu Lys Leu Lys Ser Glu Cys Phe Ser Lys Arg Thr		
610	615	620
Leu Asp His Asp Arg Cys Asp Gln His Thr Asp Cys Tyr Ser Cys Thr		
625	630	635
Ala Asn Thr Asn Asp Cys His Trp Cys Asn Asp His Cys Val Pro Arg		
645	650	655
Asn His Ser Cys Ser Glu Gly Gln Ile Ser Ile Phe Arg Tyr Glu Asn		

660

665

670

Cys Pro Lys Asp Asn Pro Met Tyr Tyr Cys Asn Lys Lys Thr Ser Cys
 675 680 685

Arg Ser Cys Ala Leu Asp Gln Asn Cys Gln Trp Glu Pro Arg Asn Gln
 690 695 700

Glu Cys Ile Ala Leu Pro Glu Asn Ile Cys Gly Ile Gly Trp His Leu
 705 710 715 720

Val Gly Asn Ser Cys Leu Lys Ile Thr Thr Ala Lys Glu Asn Tyr Asp
 725 730 735

Asn Ala Lys Leu Phe Cys Arg Asn His Asn Ala Leu Leu Ala Ser Leu
 740 745 750

Thr Thr Gln Lys Lys Val Glu Phe Val Leu Lys Gln Leu Arg Ile Met
 755 760 765

Gln Ser Ser Gln Ser Met Ser Lys Leu Thr Leu Thr Pro Trp Val Gly
 770 775 780

Leu Arg Lys Ile Asn Val Ser Tyr Trp Cys Trp Glu Asp Met Ser Pro
 785 790 795 800

Phe Thr Asn Ser Leu Leu Gln Trp Met Pro Ser Glu Pro Ser Asp Ala
 805 810 815

Gly Phe Cys Gly Ile Leu Ser Glu Pro Ser Thr Arg Gly Leu Lys Ala
 820 825 830

Ala Thr Cys Ile Asn Pro Leu Asn Gly Ser Val Cys Glu Arg Pro Ala
 835 840 845

Asn His Ser Ala Lys Gln Cys Arg Thr Pro Cys Ala Leu Arg Thr Ala
 850 855 860

Cys Gly Asp Cys Thr Ser Gly Ser Ser Glu Cys Met Trp Cys Ser Asn
 865 870 875 880

Met Lys Gln Cys Val Asp Ser Asn Ala Tyr Val Ala Ser Phe Pro Phe
 885 890 895

Gly Gln Cys Met Glu Trp Tyr Thr Met Ser Thr Cys Pro Pro Glu Asn
 900 905 910

Cys Ser Gly Tyr Cys Thr Cys Ser His Cys Leu Glu Gln Pro Gly Cys
 915 920 925

Gly Trp Cys Thr Asp Pro Ser Asn Thr Gly Lys Gly Lys Cys Ile Glu
 930 935 940

Gly Ser Tyr Lys Gly Pro Val Lys Met Pro Ser Gln Ala Pro Thr Gly
 945 950 955 960

Asn Phe Tyr Pro Gln Pro Leu Leu Asn Ser Ser Met Cys Leu Glu Asp

965

970

975

Ser Arg Tyr Asn Trp Ser Phe Ile His Cys Pro Ala Cys Gln Cys Asn
 980 985 990

Gly His Ser Lys Cys Ile Asn Gln Ser Ile Cys Glu Lys Cys Glu Asn
 995 1000 1005

Leu Thr Thr Gly Lys His Cys Glu Thr Cys Ile Ser Gly Phe Tyr Gly
 1010 1015 1020

Asp Pro Thr Asn Gly Gly Lys Cys Gln Pro Cys Lys Cys Asn Gly His
 1025 1030 1035 1040

Ala Ser Leu Cys Asn Thr Asn Thr Gly Lys Cys Phe Cys Thr Thr Lys
 1045 1050 1055

Gly Val Lys Gly Asp Glu Cys Gln Leu Cys Glu Val Glu Asn Arg Tyr
 1060 1065 1070

Gln Gly Asn Pro Leu Arg Gly Thr Cys Tyr Tyr Thr Leu Leu Ile Asp
 1075 1080 1085

Tyr Gln Phe Thr Phe Ser Leu Ser Gln Glu Asp Asp Arg Tyr Tyr Thr
 1090 1095 1100

Ala Ile Asn Phe Val Ala Thr Pro Asp Glu Gln Asn Arg Asp Leu Asp
 1105 1110 1115 1120

Met Phe Ile Asn Ala Ser Lys Asn Phe Asn Leu Asn Ile Thr Trp Ala
 1125 1130 1135

Ala Ser Phe Ser Ala Gly Thr Gln Ala Gly Glu Glu Met Pro Val Val
 1140 1145 1150

Ser Lys Thr Asn Ile Lys Glu Tyr Lys Asp Ser Phe Ser Asn Glu Lys
 1155 1160 1165

Phe Asp Phe Arg Asn His Pro Asn Ile Thr Phe Phe Val Tyr Val Ser
 1170 1175 1180

Asn Phe Thr Trp Pro Ile Lys Ile Gln Val Gln Thr Glu Gln
 1185 1190 1195

<210> 36
 <211> 1275
 <212> PRT
 <213> Rattus norvegicus

<400> 36
 Met Val Ala Ala Ala Ala Ala Glu Ala Thr Glu Ala Arg Leu Arg
 1 5 10 15

Gly Tyr Thr Thr Ala Thr Ala Ala Pro Ala Gly Trp Lys Glu Arg Gln
 20 25 30

His Arg Pro Cys Ala Ala Thr Gly Ala Trp Arg Pro Trp Pro Arg Ala
35 40 45

Gly Leu Cys Leu Pro Arg Val Leu Ser Arg Ala Leu Ser Pro Pro Pro
50 55 60

Leu Leu Pro Leu Leu Pro Leu Leu Phe Ser Leu Leu Leu Pro Leu
65 70 75 80

Pro Arg Glu Ala Glu Ala Ala Ala Val Ala Ala Ala Val Ser Gly Ser
85 90 95

Ala Ala Ala Glu Ala Lys Glu Cys Asp Arg Pro Cys Val Asn Gly Gly
100 105 110

Arg Cys Asn Pro Gly Thr Gly Gln Cys Val Cys Pro Thr Gly Trp Val
115 120 125

Gly Glu Gln Cys Gln His Cys Gly Gly Arg Phe Arg Leu Thr Gly Ser
130 135 140

Ser Gly Phe Val Thr Asp Gly Pro Gly Asn Tyr Lys Tyr Lys Thr Lys
145 150 155 160

Cys Thr Trp Leu Ile Glu Gly Gln Pro Asn Lys Ile Met Arg Leu Arg
165 170 175

Phe Asn His Phe Ala Thr Glu Cys Ser Trp Asp His Leu Tyr Val Tyr
180 185 190

Asp Gly Asp Ser Ile Tyr Ala Pro Leu Ile Ala Ala Phe Ser Gly Leu
195 200 205

Ile Val Pro Glu Arg Asp Gly Asn Glu Thr Ala Pro Glu Val Thr Val
210 215 220

Thr Ser Gly Tyr Ala Leu Leu His Phe Phe Ser Asp Ala Ala Tyr Asn
225 230 235 240

Leu Thr Gly Phe Asn Ile Thr Tyr Asn Phe Asp Met Cys Pro Asn Asn
245 250 255

Cys Ser Gly Arg Gly Glu Cys Lys Ser Ser Asn Ser Ser Thr Val
260 265 270

Glu Cys Glu Cys Ser Glu Asn Trp Lys Gly Glu Ser Cys Asp Ile Pro
275 280 285

His Cys Thr Asp Asn Cys Gly Phe Pro His Arg Gly Ile Cys Asn Ala
290 295 300

Ser Asp Thr Arg Gly Cys Ser Cys Phe Pro His Trp Gln Gly Pro Gly
305 310 315 320

Cys Ser Ile Pro Val Pro Ala Asn Gln Ser Phe Trp Thr Arg Glu Glu
325 330 335

Tyr Ser Asp Leu Lys Leu Pro Arg Ala Ser His Lys Ala Glu Val Asn
 340 345 350
 Gly Asn Ile Met Trp Val Val Gly Gly Tyr Met Phe Asn His Ser Asp
 355 360 365
 Tyr Ser Met Val Leu Ala Tyr Asp Leu Ala Ser Arg Glu Trp Leu Ser
 370 375 380
 Leu Asn His Ser Val Asn Ser Val Val Val Arg Tyr Gly His Ser Leu
 385 390 395 400
 Ala Leu His Lys Asp Lys Ile Tyr Met Tyr Gly Gly Lys Ile Asp Ser
 405 410 415
 Thr Gly Asn Val Thr Asn Glu Leu Arg Val Phe His Ile His Asn Glu
 420 425 430
 Ser Trp Val Leu Leu Thr Pro Lys Ala Lys Asp Gln Tyr Ala Val Val
 435 440 445
 Gly His Ser Ala His Ile Val Thr Leu Ser Ser Gly Arg Val Val Met
 450 455 460
 Leu Val Ile Phe Gly His Cys Pro Leu Tyr Gly Tyr Ile Ser Val Val
 465 470 475 480
 Gln Glu Tyr Asp Leu Glu Lys Asn Thr Trp Ser Ile Leu Gln Thr Gln
 485 490 495
 Gly Ala Leu Val Gln Gly Gly Tyr Gly His Ser Ser Val Tyr Asp His
 500 505 . 510
 Arg Thr Lys Ala Leu Tyr Val His Gly Gly Tyr Lys Ala Phe Ser Ala
 515 520 525
 Asn Lys Tyr Arg Leu Ala Asp Asp Leu Tyr Arg Tyr His Val Asp Thr
 530 535 540
 Gln Met Trp Thr Ile Leu Lys Asp Ser Arg Phe Phe Arg Tyr Leu His
 545 550 555 560
 Thr Ala Val Ile Val Ser Gly Thr Met Leu Val Phe Gly Gly Asn Thr
 565 570 575
 His Asn Asp Thr Ser Met Ser His Gly Ala Lys Cys Phe Ser Ser Asp
 580 585 590
 Phe Met Ala Tyr Asp Ile Ala Cys Asp Arg Trp Ser Val Leu Pro Arg
 595 600 605
 Pro Glu Leu His His Asp Val Asn Arg Phe Gly His Ser Ala Val Leu
 610 615 620
 His Asn Ser Thr Met Tyr Val Phe Gly Gly Phe Asn Ser Leu Leu Leu
 625 630 635 640

Ser Asp Val Leu Val Phe Thr Ser Glu Gln Cys Asp Ala His Arg Ser
 645 650 655
 Glu Ala Ala Cys Val Ala Ala Gly Pro Gly Ile Arg Cys Leu Trp Asp
 660 665 670
 Thr Gln Ser Ser Arg Cys Thr Ser Trp Glu Leu Ala Thr Glu Glu Gln
 675 680 685
 Ala Glu Lys Leu Lys Ser Glu Cys Phe Ser Lys Arg Thr Leu Asp His
 690 695 700
 Asp Arg Cys Asp Gln His Thr Asp Cys Tyr Ser Cys Thr Ala Asn Thr
 705 710 715 720
 Asn Asp Cys His Trp Cys Asn Asp His Cys Val Pro Val Asn His Ser
 725 730 735
 Cys Thr Glu Gly Gln Ile Ser Ile Ala Lys Tyr Asp Asn Cys Pro Lys
 740 745 750
 Asp Asn Pro Met Tyr Tyr Cys Asn Lys Lys Thr Ser Cys Arg Ser Cys
 755 760 765
 Ala Leu Asp Gln Asn Cys Gln Trp Glu Pro Arg Asn Gln Glu Cys Ile
 770 775 780
 Ala Leu Pro Glu Asn Ile Cys Gly Ile Gly Trp His Leu Val Gly Asn
 785 790 795 800
 Ser Cys Leu Lys Ile Thr Thr Ala Lys Glu Asn Tyr Asp Asn Ala Lys
 805 810 815
 Leu Ser Cys Arg Asn His Asn Ala Phe Leu Ala Ser Leu Thr Ser Gln
 820 825 830
 Lys Lys Val Glu Phe Val Leu Lys Gln Leu Arg Leu Met Gln Ser Ser
 835 840 845
 Gln Ser Thr Ser Lys Leu Thr Leu Thr Pro Trp Val Gly Leu Arg Lys
 850 855 860
 Ile Asn Val Ser Tyr Trp Cys Trp Glu Asp Met Ser Pro Phe Thr Asn
 865 870 875 880
 Ser Leu Leu Gln Trp Met Pro Ser Glu Pro Ser Asp Ala Gly Phe Cys
 885 890 895
 Gly Ile Leu Ser Glu Pro Ser Thr Arg Gly Leu Lys Ala Ala Thr Cys
 900 905 910
 Ile Asn Pro Leu Asn Gly Ser Val Cys Glu Arg Pro Ala Asn His Ser
 915 920 925
 Ala Lys Gln Cys Arg Thr Pro Cys Ala Leu Arg Thr Ala Cys Gly Glu
 930 935 940

Cys Thr Ser Ser Ser Glu Cys Met Trp Cys Ser Asn Met Lys Gln
 945 950 955 960

Cys Val Asp Ser Asn Ala Tyr Val Ala Ser Phe Pro Phe Gly Gln Cys
 965 970 975

Met Glu Trp Tyr Thr Met Ser Ser Cys Pro Pro Glu Asn Cys Ser Gly
 980 985 990

Tyr Cys Thr Cys Ser His Cys Leu Glu Gln Pro Gly Cys Gly Trp Cys
 995 1000 1005

Thr Asp Pro Ser Asn Thr Gly Lys Gly Lys Cys Ile Glu Gly Ser Tyr
 1010 1015 1020

Lys Gly Pro Val Lys Met Pro Ser His Ala Ser Thr Gly Asn Val Tyr
 1025 1030 1035 1040

Pro Gln Pro Leu Leu Asn Ser Ser Met Cys Leu Glu Asp Ser Arg Tyr
 1045 1050 1055

Asn Trp Ser Phe Ile His Cys Pro Ala Cys Gln Cys Asn Gly His Ser
 1060 1065 1070

Lys Cys Ile Asn Gln Ser Ile Cys Glu Lys Cys Glu Asp Leu Thr Thr
 1075 1080 1085

Gly Lys His Cys Glu Thr Cys Ile Ser Gly Phe Tyr Gly Asp Pro Thr
 1090 1095 1100

Asn Gly Gly Lys Cys Gln Pro Cys Lys Cys Asn Gly His Ala Ser Leu
 1105 1110 1115 1120

Cys Asn Thr Asn Thr Gly Lys Cys Phe Cys Thr Thr Lys Gly Val Lys
 1125 1130 1135

Gly Glu Glu Cys Gln Leu Cys Glu Val Glu Asn Arg Tyr Gln Gly Asn
 1140 1145 1150

Pro Leu Lys Gly Thr Cys Tyr Tyr Thr Leu Leu Ile Asp Tyr Gln Phe
 1155 1160 1165

Thr Phe Ser Leu Ser Gln Glu Asp Asp Arg Tyr Tyr Thr Ala Ile Asn
 1170 1175 1180

Phe Val Ala Thr Pro Asp Glu Gln Asn Arg Asp Leu Asp Met Phe Ile
 1185 1190 1195 1200

Asn Ala Ser Lys Asn Phe Asn Leu Asn Ile Thr Trp Ala Thr Ser Phe
 1205 1210 1215

Pro Ala Gly Thr Gln Thr Gly Glu Val Pro Val Val Ser Lys Thr
 1220 1225 1230

Asn Ile Lys Glu Tyr Lys Asp Ser Phe Ser Asn Glu Lys Phe Asp Phe
 1235 1240 1245

Arg Asn His Pro Asn Ile Thr Phe Phe Val Tyr Val Ser Asn Phe Thr
1250 1255 1260

Trp Pro Ile Lys Ile Gln Val Arg Val Thr Ser
1265 1270 1275

<210> 37
<211> 883
<212> PRT
<213> Homo sapiens

<400> 37
Met Leu Glu Met Asn Ala Arg Ser Leu Gln Gln Lys Leu Glu Thr Glu
1 5 10 15

Arg Glu Leu Lys Gln Arg Leu Leu Glu Glu Gln Ala Lys Leu Gln Gln
20 25 30

Gln Met Asp Leu Gln Lys Asn His Ile Phe Arg Leu Thr Gln Gly Leu
35 40 45

Gln Glu Ala Leu Asp Arg Ala Asp Leu Leu Lys Thr Glu Arg Ser Asp
50 55 60

Leu Glu Tyr Gln Leu Glu Asn Ile Gln Val Leu Tyr Ser His Glu Lys
65 70 75 80

Val Lys Met Glu Gly Thr Ile Ser Gln Gln Thr Lys Leu Ile Asp Phe
85 90 95

Leu Gln Ala Lys Met Asp Gln Pro Ala Lys Lys Lys Val Pro Leu
100 105 110

Gln Tyr Asn Glu Leu Lys Leu Ala Leu Glu Lys Glu Lys Ala Arg Cys
115 120 125

Ala Glu Leu Glu Ala Leu Gln Lys Thr Arg Ile Glu Leu Arg Ser
130 135 140

Ala Arg Glu Glu Ala Ala His Arg Lys Ala Thr Asp His Pro His Pro
145 150 155 160

Ser Thr Pro Ala Thr Ala Arg Gln Gln Ile Ala Met Ser Ala Ile Val
165 170 175

Arg Ser Pro Glu His Gln Pro Ser Ala Met Ser Leu Leu Ala Pro Pro
180 185 190

Ser Ser Arg Arg Lys Glu Ser Ser Thr Pro Glu Glu Phe Ser Arg Arg
195 200 205

Leu Lys Glu Arg Met His His Asn Ile Pro His Arg Phe Asn Val Gly
210 215 220

Leu Asn Met Arg Ala Thr Lys Cys Ala Val Cys Leu Asp Thr Val His
225 230 235 240

Phe Gly Arg Gln Ala Ser Lys Cys Leu Glu Cys Gln Val Met Cys His
245 250 255

Pro Lys Cys Ser Thr Cys Leu Pro Ala Thr Cys Gly Leu Pro Ala Glu
260 265 270

Tyr Ala Thr His Phe Thr Glu Ala Phe Cys Arg Asp Lys Met Asn Ser
275 280 285

Pro Gly Leu Gln Thr Lys Glu Pro Ser Ser Ser Leu His Leu Glu Gly
290 295 300

Trp Met Lys Val Pro Arg Asn Asn Lys Arg Gly Gln Gln Gly Trp Asp
305 310 315 320

Arg Lys Tyr Ile Val Leu Glu Gly Ser Lys Val Leu Ile Tyr Asp Asn
325 330 335

Glu Ala Arg Glu Ala Gly Gln Arg Pro Val Glu Glu Phe Glu Leu Cys
340 345 350

Leu Pro Asp Gly Asp Val Ser Ile His Gly Ala Val Gly Ala Ser Glu
355 360 365

Leu Ala Asn Thr Ala Lys Ala Asp Val Pro Tyr Ile Leu Lys Met Glu
370 375 380

Ser His Pro His Thr Thr Cys Trp Pro Gly Arg Thr Leu Tyr Leu Leu
385 390 395 400

Ala Pro Ser Phe Pro Asp Lys Gln Arg Trp Val Thr Ala Leu Glu Ser
405 410 415

Val Val Ala Gly Gly Arg Val Ser Arg Glu Lys Ala Glu Ala Asp Ala
420 425 430

Lys Leu Leu Gly Asn Ser Leu Leu Lys Leu Glu Gly Asp Asp Arg Leu
435 440 445

Asp Met Asn Cys Thr Leu Pro Phe Ser Asp Gln Val Val Leu Val Gly
450 455 460

Thr Glu Glu Gly Leu Tyr Ala Leu Asn Val Leu Lys Asn Ser Leu Thr
465 470 475 480

His Val Pro Gly Ile Gly Ala Val Phe Gln Ile Tyr Ile Ile Lys Asp
485 490 495

Leu Glu Lys Leu Leu Met Ile Ala Gly Glu Glu Arg Ala Leu Cys Leu
500 505 510

Val Asp Val Lys Lys Val Lys Gln Ser Leu Ala Gln Ser His Leu Pro
515 520 525

Ala Gln Pro Asp Ile Ser Pro Asn Ile Phe Glu Ala Val Lys Gly Cys
530 535 540

His Leu Phe Gly Ala Gly Lys Ile Glu Asn Gly Leu Cys Ile Cys Ala
 545 550 555 560
 Ala Met Pro Ser Lys Val Val Ile Leu Arg Tyr Asn Glu Asn Leu Ser
 565 570 575
 Lys Tyr Cys Ile Arg Lys Glu Ile Glu Thr Ser Glu Pro Cys Ser Cys
 580 585 590
 Ile His Phe Thr Asn Tyr Ser Ile Leu Ile Gly Thr Asn Lys Phe Tyr
 595 600 605
 Glu Ile Asp Met Lys Gln Tyr Thr Leu Glu Glu Phe Leu Asp Lys Asn
 610 615 620
 Asp His Ser Leu Ala Pro Ala Val Phe Ala Ala Ser Ser Asn Ser Phe
 625 630 635 640
 Pro Val Ser Ile Val Gln Val Asn Ser Ala Gly Gln Arg Glu Glu Tyr
 645 650 655
 Leu Leu Cys Phe His Glu Phe Gly Val Phe Val Asp Ser Tyr Gly Arg
 660 665 670
 Arg Ser Arg Thr Asp Asp Leu Lys Trp Ser Arg Leu Pro Leu Ala Phe
 675 680 685
 Ala Tyr Arg Glu Pro Tyr Leu Phe Val Thr His Phe Asn Ser Leu Glu
 690 695 700
 Val Ile Glu Ile Gln Ala Arg Ser Ser Ala Gly Thr Pro Ala Arg Ala
 705 710 715 720
 Tyr Leu Asp Ile Pro Asn Pro Arg Tyr Leu Gly Pro Ala Ile Ser Ser
 725 730 735
 Gly Ala Ile Tyr Leu Ala Ser Ser Tyr Gln Asp Lys Leu Arg Val Ile
 740 745 750
 Cys Cys Lys Gly Asn Leu Val Lys Glu Ser Gly Thr Glu His His Arg
 755 760 765
 Gly Pro Ser Thr Ser Arg Ser Ser Pro Asn Lys Arg Gly Pro Pro Thr
 770 775 780
 Tyr Asn Glu His Ile Thr Lys Arg Val Ala Ser Ser Pro Ala Pro Pro
 785 790 795 800
 Glu Gly Pro Ser His Pro Arg Glu Pro Ser Thr Pro His Arg Tyr Arg
 805 810 815
 Glu Gly Arg Thr Glu Leu Arg Arg Asp Lys Ser Pro Gly Arg Pro Leu
 820 825 830
 Glu Arg Glu Lys Ser Pro Gly Arg Met Leu Ser Thr Arg Arg Glu Arg
 835 840 845

Ser Pro Gly Arg Leu Phe Glu Asp Ser Ser Arg Gly Arg Leu Pro Ala
850 855 860

Gly Ala Val Arg Thr Pro Leu Ser Gln Val Asn Lys Val Trp Asp Gln
865 870 875 880

Ser Ser Val

<210> 38
<211> 1286
<212> PRT
<213> Homo sapiens

<400> 38
Val Leu Asp Asn Gln Ile Lys Lys Asp Leu Ala Asp Lys Glu Thr Leu
1 5 10 15

Glu Asn Met Met Gln Arg His Glu Glu Glu Ala His Glu Lys Gly Lys
20 25 30

Ile Leu Ser Glu Gln Lys Ala Met Ile Asn Ala Met Asp Ser Lys Ile
35 40 45

Arg Ser Leu Glu Gln Arg Ile Val Glu Leu Ser Glu Ala Asn Lys Leu
50 55 60

Ala Ala Asn Ser Ser Leu Phe Thr Gln Arg Asn Met Lys Ala Gln Glu
65 70 75 80

Glu Met Ile Ser Glu Leu Arg Gln Gln Lys Phe Tyr Leu Glu Thr Gln
85 90 95

Ala Gly Lys Leu Glu Ala Gln Asn Arg Lys Leu Glu Glu Gln Leu Glu
100 105 110

Lys Ile Ser His Gln Asp His Ser Asp Lys Asn Arg Leu Leu Glu Leu
115 120 125

Glu Thr Arg Leu Arg Glu Val Ser Leu Glu His Glu Glu Gln Lys Leu
130 135 140

Glu Leu Lys Arg Gln Leu Thr Glu Leu Gln Leu Ser Leu Gln Glu Arg
145 150 155 160

Glu Ser Gln Leu Thr Ala Leu Gln Ala Ala Arg Ala Ala Leu Glu Ser
165 170 175

Gln Leu Arg Gln Ala Lys Thr Glu Leu Glu Glu Thr Thr Ala Glu Ala
180 185 190

Glu Glu Glu Ile Gln Ala Leu Thr Ala His Arg Asp Glu Ile Gln Arg
195 200 205

Lys Phe Asp Ala Leu Arg Asn Ser Cys Thr Val Ile Thr Asp Leu Glu

210 215 220
Glu Gln Leu Asn Gln Leu Thr Glu Asp Asn Ala Glu Leu Asn Asn Gln
225 230 235 240
Asn Phe Tyr Leu Ser Lys Gln Leu Asp Glu Ala Ser Gly Ala Asn Asp
245 250 255
Glu Ile Val Gln Leu Arg Ser Glu Val Asp His Leu Arg Arg Glu Ile
260 265 270
Thr Glu Arg Glu Met Gln Leu Thr Ser Gln Lys Gln Thr Met Glu Ala
275 280 285
Leu Lys Thr Thr Cys Thr Met Leu Glu Glu Gln Val Met Asp Leu Glu
290 295 300
Ala Leu Asn Asp Glu Leu Leu Glu Lys Glu Arg Gln Trp Glu Ala Trp
305 310 315 320
Arg Ser Val Leu Gly Asp Glu Lys Ser Gln Phe Glu Cys Arg Val Arg
325 330 335
Glu Leu Gln Arg Met Leu Asp Thr Glu Lys Gln Ser Arg Ala Arg Ala
340 345 350
Asp Gln Arg Ile Thr Glu Ser Arg Gln Val Val Glu Leu Ala Val Lys
355 360 365
Glu His Lys Ala Glu Ile Leu Ala Leu Gln Gln Ala Leu Lys Glu Gln
370 375 380
Lys Leu Lys Ala Glu Ser Leu Ser Asp Lys Leu Asn Asp Leu Glu Lys
385 390 395 400
Lys His Ala Met Leu Glu Met Asn Ala Arg Ser Leu Gln Gln Lys Leu
405 410 415
Glu Thr Glu Arg Glu Leu Lys Gln Arg Leu Leu Glu Glu Gln Ala Lys
420 425 430
Leu Gln Gln Gln Met Asp Leu Gln Lys Asn His Ile Phe Arg Leu Thr
435 440 445
Gln Gly Leu Gln Glu Ala Leu Asp Arg Ala Asp Leu Leu Lys Thr Glu
450 455 460
Arg Ser Asp Leu Glu Tyr Gln Leu Glu Asn Ile Gln Val Leu Tyr Ser
465 470 475 480
His Glu Lys Val Lys Met Glu Gly Thr Ile Ser Gln Gln Thr Lys Leu
485 490 495
Ile Asp Phe Leu Gln Ala Lys Met Asp Gln Pro Ala Lys Lys Lys Lys
500 505 510
Val Pro Leu Gln Tyr Asn Glu Leu Lys Leu Ala Leu Glu Lys Glu Lys

515	520	525
Ala Arg Cys Ala Glu Leu Glu Glu Ala Leu Gln Lys Thr Arg Ile Glu		
530	535	540
Leu Arg Ser Ala Arg Glu Glu Ala Ala His Arg Lys Ala Thr Asp His		
545	550	555
Pro His Pro Ser Thr Pro Ala Thr Ala Arg Gln Gln Ile Ala Met Ser		
565	570	575
Ala Ile Val Arg Ser Pro Glu His Gln Pro Ser Ala Met Ser Leu Leu		
580	585	590
Ala Pro Pro Ser Ser Arg Arg Lys Glu Ser Ser Thr Pro Glu Glu Phe		
595	600	605
Ser Arg Arg Leu Lys Glu Arg Met His His Asn Ile Pro His Arg Phe		
610	615	620
Asn Val Gly Leu Asn Met Arg Ala Thr Lys Cys Ala Val Cys Leu Asp		
625	630	635
Thr Val His Phe Gly Arg Gln Ala Ser Lys Cys Leu Glu Cys Gln Val		
645	650	655
Met Cys His Pro Lys Cys Ser Thr Cys Leu Pro Ala Thr Cys Gly Leu		
660	665	670
Pro Ala Glu Tyr Ala Thr His Phe Thr Glu Ala Phe Cys Arg Asp Lys		
675	680	685
Met Asn Ser Pro Gly Leu Gln Thr Lys Glu Pro Ser Ser Ser Leu His		
690	695	700
Leu Glu Gly Trp Met Lys Val Pro Arg Asn Asn Lys Arg Gly Gln Gln		
705	710	715
Gly Trp Asp Arg Lys Tyr Ile Val Leu Glu Gly Ser Lys Val Leu Ile		
725	730	735
Tyr Asp Asn Glu Ala Arg Glu Ala Gly Gln Arg Pro Val Glu Glu Phe		
740	745	750
Glu Leu Cys Leu Pro Asp Gly Asp Val Ser Ile His Gly Ala Val Gly		
755	760	765
Ala Ser Glu Leu Ala Asn Thr Ala Lys Ala Asp Val Pro Tyr Ile Leu		
770	775	780
Lys Met Glu Ser His Pro His Thr Thr Cys Trp Pro Gly Arg Thr Leu		
785	790	795
Tyr Leu Leu Ala Pro Ser Phe Pro Asp Lys Gln Arg Trp Val Thr Ala		
805	810	815
Leu Glu Ser Val Val Ala Gly Gly Arg Val Ser Arg Glu Lys Ala Glu		

820	825	830
Ala Asp Ala Lys Leu Leu Gly Asn Ser Leu Leu Lys Leu Glu Gly Asp		
835	840	845
Asp Arg Leu Asp Met Asn Cys Thr Leu Pro Phe Ser Asp Gln Val Val		
850	855	860
Leu Val Gly Thr Glu Glu Gly Leu Tyr Ala Leu Asn Val Leu Lys Asn		
865	870	875
880		
Ser Leu Thr His Val Pro Gly Ile Gly Ala Val Phe Gln Ile Tyr Ile		
885	890	895
Ile Lys Asp Leu Glu Lys Leu Leu Met Ile Ala Gly Glu Glu Arg Ala		
900	905	910
Leu Cys Leu Val Asp Val Lys Lys Val Lys Gln Ser Leu Ala Gln Ser		
915	920	925
His Leu Pro Ala Gln Pro Asp Ile Ser Pro Asn Ile Phe Glu Ala Val		
930	935	940
Lys Gly Cys His Leu Phe Gly Ala Gly Lys Ile Glu Asn Gly Leu Cys		
945	950	955
960		
Ile Cys Ala Ala Met Pro Ser Lys Val Val Ile Leu Arg Tyr Asn Glu		
965	970	975
Asn Leu Ser Lys Tyr Cys Ile Arg Lys Glu Ile Glu Thr Ser Glu Pro		
980	985	990
Cys Ser Cys Ile His Phe Thr Asn Tyr Ser Ile Leu Ile Gly Thr Asn		
995	1000	1005
Lys Phe Tyr Glu Ile Asp Met Lys Gln Tyr Thr Leu Glu Glu Phe Leu		
1010	1015	1020
Asp Lys Asn Asp His Ser Leu Ala Pro Ala Val Phe Ala Ala Ser Ser		
1025	1030	1035
1040		
Asn Ser Phe Pro Val Ser Ile Val Gln Val Asn Ser Ala Gly Gln Arg		
1045	1050	1055
Glu Glu Tyr Leu Leu Cys Phe His Glu Phe Gly Val Phe Val Asp Ser		
1060	1065	1070
Tyr Gly Arg Arg Ser Arg Thr Asp Asp Leu Lys Trp Ser Arg Leu Pro		
1075	1080	1085
Leu Ala Phe Ala Tyr Arg Glu Pro Tyr Leu Phe Val Thr His Phe Asn		
1090	1095	1100
Ser Leu Glu Val Ile Glu Ile Gln Ala Arg Ser Ser Ala Gly Thr Pro		
1105	1110	1115
1120		
Ala Arg Ala Tyr Leu Asp Ile Pro Asn Pro Arg Tyr Leu Gly Pro Ala		

1125

1130

1135

Ile Ser Ser Gly Ala Ile Tyr Leu Ala Ser Ser Tyr Gln Asp Lys Leu
1140 1145 1150

Arg Val Ile Cys Cys Lys Gly Asn Leu Val Lys Glu Ser Gly Thr Glu
1155 1160 1165

His His Arg Gly Pro Ser Thr Ser Arg Ser Ser Pro Asn Lys Arg Gly
1170 1175 1180

Pro Pro Thr Tyr Asn Glu His Ile Thr Lys Arg Val Ala Ser Ser Pro
1185 1190 1195 1200

Ala Pro Pro Glu Gly Pro Ser His Pro Arg Glu Pro Ser Thr Pro His
1205 1210 1215

Arg Tyr Arg Glu Gly Arg Thr Glu Leu Arg Arg Asp Lys Ser Pro Gly
1220 1225 1230

Arg Pro Leu Glu Arg Glu Lys Ser Pro Gly Arg Met Leu Ser Thr Arg
1235 1240 1245

Arg Glu Arg Ser Pro Gly Arg Leu Phe Glu Asp Ser Ser Arg Gly Arg
1250 1255 1260

Leu Pro Ala Gly Ala Val Arg Thr Pro Leu Ser Gln Val Asn Lys Val
1265 1270 1275 1280

Trp Asp Gln Ser Ser Val
1285

<210> 39
<211> 940
<212> PRT
<213> Homo sapiens

<400> 39
Gln Ser Arg Ala Arg Ala Asp Gln Arg Ile Thr Glu Ser Arg Gln Val
1 5 10 15

Val Glu Leu Ala Val Lys Glu His Lys Ala Glu Ile Leu Ala Leu Gln
20 25 30

Gln Ala Leu Lys Glu Gln Lys Leu Lys Ala Glu Ser Leu Ser Asp Lys
35 40 45

Leu Asn Asp Leu Glu Lys Lys His Ala Met Leu Glu Met Asn Ala Arg
50 55 60 80

Ser Leu Gln Gln Lys Leu Glu Thr Glu Arg Glu Leu Lys Gln Arg Leu
65 70 75 80

Leu Glu Glu Gln Ala Lys Leu Gln Gln Met Asp Leu Gln Lys Asn
85 90 95

His	Ile	Phe	Arg	Leu	Thr	Gln	Gly	Leu	Gln	Glu	Ala	Leu	Asp	Arg	Ala
100								105					110		
Asp	Leu	Leu	Lys	Thr	Glu	Arg	Ser	Asp	Leu	Glu	Tyr	Gln	Leu	Glu	Asn
115								120					125		
Ile	Gln	Val	Leu	Tyr	Ser	His	Glu	Lys	Val	Lys	Met	Glu	Gly	Thr	Ile
130								135				140			
Ser	Gln	Gln	Thr	Lys	Leu	Ile	Asp	Phe	Leu	Gln	Ala	Lys	Met	Asp	Gln
145								150				155			160
Pro	Ala	Lys	Lys	Lys	Val	Pro	Leu	Gln	Tyr	Asn	Glu	Leu	Lys	Leu	
								165				170			175
Ala	Leu	Glu	Lys	Glu	Lys	Ala	Arg	Cys	Ala	Glu	Leu	Glu	Glu	Ala	Leu
								180				185			190
Gln	Lys	Thr	Arg	Ile	Glu	Leu	Arg	Ser	Ala	Arg	Glu	Glu	Ala	Ala	His
								195				200			205
Arg	Lys	Ala	Thr	Asp	His	Pro	His	Pro	Ser	Thr	Pro	Ala	Thr	Ala	Arg
								210				215			220
Gln	Gln	Ile	Ala	Met	Ser	Ala	Ile	Val	Arg	Ser	Pro	Glu	His	Gln	Pro
								225				230			240
Ser	Ala	Met	Ser	Leu	Leu	Ala	Pro	Pro	Ser	Ser	Arg	Arg	Lys	Glu	Ser
								245				250			255
Ser	Thr	Pro	Glu	Glu	Phe	Ser	Arg	Arg	Leu	Lys	Glu	Arg	Met	His	His
								260				265			270
Asn	Ile	Pro	His	Arg	Phe	Asn	Val	Gly	Leu	Asn	Met	Arg	Ala	Thr	Lys
								275				280			285
Cys	Ala	Val	Cys	Leu	Asp	Thr	Val	His	Phe	Gly	Arg	Gln	Ala	Ser	Lys
								290				295			300
Cys	Leu	Glu	Cys	Gln	Val	Met	Cys	His	Pro	Lys	Cys	Ser	Thr	Cys	Leu
								305				310			320
Pro	Ala	Thr	Cys	Gly	Leu	Pro	Ala	Glu	Tyr	Ala	Thr	His	Phe	Thr	Glu
								325				330			335
Ala	Phe	Cys	Arg	Asp	Lys	Met	Asn	Ser	Pro	Gly	Leu	Gln	Thr	Lys	Glu
								340				345			350
Pro	Ser	Ser	Ser	Leu	His	Leu	Glu	Gly	Trp	Met	Lys	Val	Pro	Arg	Asn
								355				360			365
Asn	Lys	Arg	Gly	Gln	Gln	Gly	Trp	Asp	Arg	Lys	Tyr	Ile	Val	Leu	Glu
								370				375			380
Gly	Ser	Lys	Val	Leu	Ile	Tyr	Asp	Asn	Glu	Ala	Arg	Glu	Ala	Gly	Gln
								385				390			400

Arg Pro Val Glu Glu Phe Glu Leu Cys Leu Pro Asp Gly Asp Val Ser
 405 410 415

 Ile His Gly Ala Val Gly Ala Ser Glu Leu Ala Asn Thr Ala Lys Ala
 420 425 430

 Asp Val Pro Tyr Ile Leu Lys Met Glu Ser His Pro His Thr Thr Cys
 435 440 445

 Trp Pro Gly Arg Thr Leu Tyr Leu Leu Ala Pro Ser Phe Pro Asp Lys
 450 455 460

 Gln Arg Trp Val Thr Ala Leu Glu Ser Val Val Ala Gly Gly Arg Val
 465 470 475 480

 Ser Arg Glu Lys Ala Glu Ala Asp Ala Lys Leu Leu Gly Asn Ser Leu
 485 490 495

 Leu Lys Leu Glu Gly Asp Asp Arg Leu Asp Met Asn Cys Thr Leu Pro
 500 505 510

 Phe Ser Asp Gln Val Val Leu Val Gly Thr Glu Glu Gly Leu Tyr Ala
 515 520 525

 Leu Asn Val Leu Lys Asn Ser Leu Thr His Val Pro Gly Ile Gly Ala
 530 535 540

 Val Phe Gln Ile Tyr Ile Ile Lys Asp Leu Glu Lys Leu Leu Met Ile
 545 550 555 560

 Ala Gly Glu Glu Arg Ala Leu Cys Leu Val Asp Val Lys Lys Val Lys
 565 570 575

 Gln Ser Leu Ala Gln Ser His Leu Pro Ala Gln Pro Asp Ile Ser Pro
 580 585 590

 Asn Ile Phe Glu Ala Val Lys Gly Cys His Leu Phe Gly Ala Gly Lys
 595 600 605

 Ile Glu Asn Gly Leu Cys Ile Cys Ala Ala Met Pro Ser Lys Val Val
 610 615 620

 Ile Leu Arg Tyr Asn Glu Asn Leu Ser Lys Tyr Cys Ile Arg Lys Glu
 625 630 635 640

 Ile Glu Thr Ser Glu Pro Cys Ser Cys Ile His Phe Thr Asn Tyr Ser
 645 650 655

 Ile Leu Ile Gly Thr Asn Lys Phe Tyr Glu Ile Asp Met Lys Gln Tyr
 660 665 670

 Thr Leu Glu Glu Phe Leu Asp Lys Asn Asp His Ser Leu Ala Pro Ala
 675 680 685

 Val Phe Ala Ala Ser Ser Asn Ser Phe Pro Val Ser Ile Val Gln Val
 690 695 700

Asn	Ser	Ala	Gly	Gln	Arg	Glu	Glu	Tyr	Leu	Leu	Cys	Phe	His	Glu	Phe
705				710					715				720		
Gly	Val	Phe	Val	Asp	Ser	Tyr	Gly	Arg	Arg	Ser	Arg	Thr	Asp	Asp	Leu
		725					730					735			
Lys	Trp	Ser	Arg	Leu	Pro	Leu	Ala	Phe	Ala	Tyr	Arg	Glu	Pro	Tyr	Leu
		740				745			750						
Phe	Val	Thr	His	Phe	Asn	Ser	Leu	Glu	Val	Ile	Glu	Ile	Gln	Ala	Arg
		755				760				765					
Ser	Ser	Ala	Gly	Thr	Pro	Ala	Arg	Ala	Tyr	Leu	Asp	Ile	Pro	Asn	Pro
		770				775				780					
Arg	Tyr	Leu	Gly	Pro	Ala	Ile	Ser	Ser	Gly	Ala	Ile	Tyr	Leu	Ala	Ser
		785				790			795			800			
Ser	Tyr	Gln	Asp	Lys	Leu	Arg	Val	Ile	Cys	Cys	Lys	Gly	Asn	Leu	Val
			805				810				815				
Lys	Glu	Ser	Gly	Thr	Glu	His	His	Arg	Gly	Pro	Ser	Thr	Ser	Arg	Ser
			820			825			830						
Ser	Pro	Asn	Lys	Arg	Gly	Pro	Pro	Thr	Tyr	Asn	Glu	His	Ile	Thr	Lys
			835			840				845					
Arg	Val	Ala	Ser	Ser	Pro	Ala	Pro	Pro	Glu	Gly	Pro	Ser	His	Pro	Arg
		850				855			860						
Glu	Pro	Ser	Thr	Pro	His	Arg	Tyr	Arg	Glu	Gly	Arg	Thr	Glu	Leu	Arg
		865				870			875			880			
Arg	Asp	Lys	Ser	Pro	Gly	Arg	Pro	Leu	Glu	Arg	Glu	Lys	Ser	Pro	Gly
			885				890			895					
Arg	Met	Leu	Ser	Thr	Arg	Arg	Glu	Arg	Ser	Pro	Gly	Arg	Leu	Phe	Glu
			900			905			910						
Asp	Ser	Ser	Arg	Gly	Arg	Leu	Pro	Ala	Gly	Ala	Val	Arg	Thr	Pro	Leu
			915			920			925						
Ser	Gln	Val	Asn	Lys	Val	Trp	Asp	Gln	Ser	Ser	Val				
			930			935			940						
<210>	40														
<211>	1641														
<212>	PRT														
<213>	Mus musculus														
<400>	40														
Pro	Phe	Val	Pro	Thr	Leu	Lys	Ser	Asp	Asp	Asp	Thr	Ser	Asn	Phe	Asp
	1				5				10			15			
Glu	Pro	Glu	Lys	Asn	Ser	Trp	Val	Ser	Ser	Ser	Val	Cys	Gln	Leu	Ser
					20			25			30				

Pro Ser Gly Phe Ser Gly Glu Glu Leu Pro Phe Val Gly Phe Ser Tyr
35 40 45

Ser Lys Ala Leu Gly Tyr Leu Gly Arg Ser Glu Ser Val Val Ser Ser
50 55 60

Leu Asp Ser Pro Ala Lys Val Ser Ser Met Glu Lys Lys Leu Leu Ile
65 70 75 80

Lys Ser Lys Glu Leu Gln Asp Ser Gln Asp Lys Cys His Lys Met Glu
85 90 95

Gln Glu Met Thr Arg Leu His Arg Arg Val Ser Glu Val Glu Ala Val
100 105 110

Leu Ser Gln Lys Glu Val Glu Leu Lys Ala Ser Glu Thr Gln Arg Ser
115 120 125

Leu Leu Glu Gln Asp Leu Ala Thr Tyr Ile Thr Glu Cys Ser Ser Leu
130 135 140

Lys Arg Ser Leu Glu Gln Ala Arg Met Glu Val Ser Gln Glu Asp Asp
145 150 155 160

Lys Ala Leu Gln Leu Leu His Asp Ile Arg Glu Gln Ser Arg Lys Leu
165 170 175

Gln Glu Ile Lys Glu Gln Glu Tyr Gln Ala Gln Val Glu Glu Met Arg
180 185 190

Leu Met Met Asn Gln Leu Glu Glu Asp Leu Val Ser Ala Arg Arg Arg
195 200 205

Ser Asp Leu Tyr Glu Ser Glu Leu Arg Glu Ser Arg Leu Ala Ala Glu
210 215 220

Glu Phe Lys Arg Lys Ala Asn Glu Cys Gln His Lys Leu Met Lys Ala
225 230 235 240

Lys Asp Gln Gly Lys Pro Glu Val Gly Glu Tyr Ser Lys Leu Glu Lys
245 250 255

Ile Asn Ala Glu Gln Gln Leu Lys Ile Gln Glu Leu Gln Glu Lys Leu
260 265 270

Glu Lys Ala Val Lys Ala Ser Thr Glu Ala Thr Glu Leu Leu Gln Asn
275 280 285

Ile Arg Gln Ala Lys Glu Arg Ala Glu Arg Glu Leu Glu Lys Leu His
290 295 300

Asn Arg Glu Asp Ser Ser Glu Gly Ile Lys Lys Lys Leu Val Glu Ala
305 310 315 320

Glu Glu Leu Glu Glu Lys His Arg Glu Ala Gln Val Ser Ala Gln His
325 330 335

Leu Glu Val His Leu Lys Gln Lys Glu Gln His Tyr Glu Glu Lys Ile
340 345 350

Lys Val Leu Asp Asn Gln Ile Lys Lys Asp Leu Ala Asp Lys Glu Ser
355 360 365

Leu Glu Asn Met Met Gln Arg His Glu Glu Glu Ala His Glu Lys Gly
370 375 380

Lys Ile Leu Ser Glu Gln Lys Ala Met Ile Asn Ala Met Asp Ser Lys
385 390 395 400

Ile Arg Ser Leu Glu Gln Arg Ile Val Glu Leu Ser Glu Ala Asn Lys
405 410 415

Leu Ala Ala Asn Ser Ser Leu Phe Thr Gln Arg Asn Met Lys Ala Gln
420 425 430

Glu Glu Met Ile Ser Glu Leu Arg Gln Gln Lys Phe Tyr Leu Glu Thr
435 440 445

Gln Ala Gly Lys Leu Glu Ala Gln Asn Arg Lys Leu Glu Glu Gln Leu
450 455 460

Glu Lys Ile Ser His Gln Asp His Ser Asp Lys Ser Arg Leu Leu Glu
465 470 475 480

Leu Glu Thr Arg Leu Arg Glu Val Ser Leu Glu His Glu Glu Gln Lys
485 490 495

Leu Glu Leu Lys Arg Gln Leu Thr Glu Leu Gln Leu Ser Leu Gln Glu
500 505 510

Arg Glu Ser Gln Leu Thr Ala Leu Gln Ala Ala Arg Ala Ala Leu Glu
515 520 525

Ser Gln Leu Arg Gln Ala Lys Thr Glu Leu Glu Glu Thr Thr Ala Glu
530 535 540

Ala Glu Glu Glu Ile Gln Ala Leu Thr Ala His Arg Asp Glu Ile Gln
545 550 555 560

Arg Lys Phe Asp Ala Leu Arg Asn Ser Cys Thr Val Ile Thr Asp Leu
565 570 575

Glu Glu Gln Leu Asn Gln Leu Thr Glu Asp Asn Ala Glu Leu Asn Asn
580 585 590

Gln Asn Phe Tyr Leu Ser Lys Gln Leu Asp Glu Ala Ser Gly Ala Asn
595 600 605

Asp Glu Ile Val Gln Leu Arg Ser Glu Val Asp His Leu Arg Arg Glu
610 615 620

Ile Thr Glu Arg Glu Met Gln Leu Thr Ser Gln Lys Gln Thr Met Glu
625 630 635 640

Ala Leu Lys Thr Thr Cys Thr Met Leu Glu Glu Gln Val Leu Asp Leu
645 650 655

Glu Ala Leu Asn Asp Glu Leu Leu Glu Lys Glu Arg Gln Trp Glu Ala
660 665 670

Trp Arg Ser Val Leu Gly Asp Glu Lys Ser Gln Phe Glu Cys Arg Val
675 680 685

Arg Glu Leu Gln Arg Met Leu Asp Thr Glu Lys Gln Ser Arg Ala Arg
690 695 700

Ala Asp Gln Arg Ile Thr Glu Ser Arg Gln Val Val Glu Leu Ala Val
705 710 715 720

Lys Glu His Lys Ala Glu Ile Leu Ala Leu Gln Gln Ala Leu Lys Glu
725 730 735

Gln Lys Leu Lys Ala Glu Ser Leu Ser Asp Lys Leu Asn Asp Leu Glu
740 745 750

Lys Lys His Ala Met Leu Glu Met Asn Ala Arg Ser Leu Gln Gln Lys
755 760 765

Leu Glu Thr Glu Arg Glu Leu Lys Gln Arg Leu Leu Glu Glu Gln Ala
770 775 780

Lys Leu Gln Gln Gln Met Asp Leu Gln Lys Asn His Ile Phe Arg Leu
785 790 795 800

Thr Gln Gly Leu Gln Glu Ala Leu Asp Arg Ala Asp Leu Leu Lys Thr
805 810 815

Glu Arg Ser Asp Leu Glu Tyr Gln Leu Glu Asn Ile Gln Val Leu Tyr
820 825 830

Ser His Glu Lys Val Lys Met Glu Gly Thr Ile Ser Gln Gln Thr Lys
835 840 845

Leu Ile Asp Phe Leu Gln Ala Lys Met Asp Gln Pro Ala Lys Lys Lys
850 855 860

Lys Val Pro Leu Gln Tyr Asn Glu Leu Lys Leu Ala Leu Glu Lys Glu
865 870 875 880

Lys Ala Arg Cys Ala Glu Leu Glu Ala Leu Gln Lys Thr Arg Ile
885 890 895

Glu Leu Arg Ser Ala Arg Glu Glu Ala Ala His Arg Lys Ala Thr Asp
900 905 910

His Pro His Pro Ser Thr Pro Ala Thr Ala Arg Gln Gln Ile Ala Met
915 920 925

Ser Ala Ile Val Arg Ser Pro Glu His Gln Pro Ser Ala Met Ser Leu
930 935 940

Leu Ala Pro Pro Ser Ser Arg Arg Lys Glu Ser Ser Thr Pro Glu Glu
 945 950 955 960

 Phe Ser Arg Arg Leu Lys Glu Arg Met His His Asn Ile Pro His Arg
 965 970 975

 Phe Asn Val Gly Leu Asn Met Arg Ala Thr Lys Cys Ala Val Cys Leu
 980 985 990

 Asp Thr Val His Phe Gly Arg Gln Ala Ser Lys Cys Leu Glu Cys Gln
 995 1000 1005

 Val Met Cys His Pro Lys Cys Ser Thr Cys Leu Pro Ala Thr Cys Gly
 1010 1015 1020

 Leu Pro Ala Glu Tyr Ala Thr His Phe Thr Glu Ala Phe Cys Arg Asp
 1025 1030 1035 1040

 Lys Met Asn Ser Pro Gly Leu Gln Ser Lys Glu Pro Gly Ser Ser Leu
 1045 1050 1055

 His Leu Glu Gly Trp Met Lys Val Pro Arg Asn Asn Lys Arg Gly Gln
 1060 1065 1070

 Gln Gly Trp Asp Arg Lys Tyr Ile Val Leu Glu Gly Ser Lys Val Leu
 1075 1080 1085

 Ile Tyr Asp Asn Glu Ala Arg Glu Ala Gly Gln Arg Pro Val Glu Glu
 1090 1095 1100

 Phe Glu Leu Cys Leu Pro Asp Gly Asp Val Ser Ile His Gly Ala Val
 1105 1110 1115 1120

 Gly Ala Ser Glu Leu Ala Asn Thr Ala Lys Ala Asp Val Pro Tyr Ile
 1125 1130 1135

 Leu Lys Met Glu Ser His Pro His Thr Thr Cys Trp Pro Gly Arg Thr
 1140 1145 1150

 Leu Tyr Leu Leu Ala Pro Ser Phe Pro Asp Lys Gln Arg Trp Val Thr
 1155 1160 1165

 Ala Leu Glu Ser Val Val Ala Gly Gly Arg Val Ser Arg Glu Lys Ala
 1170 1175 1180

 Glu Ala Asp Ala Lys Leu Leu Gly Asn Ser Leu Leu Lys Leu Glu Gly
 1185 1190 1195 1200

 Asp Asp Arg Leu Asp Met Asn Cys Thr Leu Pro Phe Ser Asp Gln Val
 1205 1210 1215

 Val Leu Val Gly Thr Glu Glu Gly Leu Tyr Ala Leu Asn Val Leu Lys
 1220 1225 1230

 Asn Ser Leu Thr His Ile Pro Gly Ile Gly Ala Val Phe Gln Ile Tyr
 1235 1240 1245

Ile Ile Lys Asp Leu Glu Lys Leu Leu Met Ile Ala Gly Glu Glu Arg
 1250 1255 1260

 Ala Leu Cys Leu Val Asp Val Lys Lys Val Lys Gln Ser Leu Ala Gln
 1265 1270 1275 1280

 Ser His Leu Pro Ala Gln Pro Asp Val Ser Pro Asn Ile Phe Glu Ala
 1285 1290 1295

 Val Lys Gly Cys His Leu Phe Ala Ala Gly Lys Ile Glu Asn Ser Leu
 1300 1305 1310

 Cys Ile Cys Ala Ala Met Pro Ser Lys Val Val Ile Leu Arg Tyr Asn
 1315 1320 1325

 Asp Asn Leu Ser Lys Tyr Cys Ile Arg Lys Glu Ile Glu Thr Ser Glu
 1330 1335 1340

 Pro Cys Ser Cys Ile His Phe Thr Asn Tyr Ser Ile Leu Ile Gly Thr
 1345 1350 1355 1360

 Asn Lys Phe Tyr Glu Ile Asp Met Lys Gln Tyr Thr Leu Asp Glu Phe
 1365 1370 1375

 Leu Asp Lys Asn Asp His Ser Leu Ala Pro Ala Val Phe Ala Ser Ser
 1380 1385 1390

 Ser Asn Ser Phe Pro Val Ser Ile Val Gln Ala Asn Ser Ala Gly Gln
 1395 1400 1405

 Arg Glu Glu Tyr Leu Leu Cys Phe His Glu Phe Gly Val Phe Val Asp
 1410 1415 1420

 Ser Tyr Gly Arg Arg Ser Arg Thr Asp Asp Leu Lys Trp Ser Arg Leu
 1425 1430 1435 1440

 Pro Leu Ala Phe Ala Tyr Arg Glu Pro Tyr Leu Phe Val Thr His Phe
 1445 1450 1455

 Asn Ser Leu Glu Val Ile Glu Ile Gln Ala Arg Ser Ser Leu Gly Ser
 1460 1465 1470

 Pro Ala Arg Ala Tyr Leu Glu Ile Pro Asn Pro Arg Tyr Leu Gly Pro
 1475 1480 1485

 Ala Ile Ser Ser Gly Ala Ile Tyr Leu Ala Ser Ser Tyr Gln Asp Lys
 1490 1495 1500

 Leu Arg Val Ile Cys Cys Lys Gly Asn Leu Val Lys Glu Ser Gly Thr
 1505 1510 1515 1520

 Glu Gln His Arg Val Pro Ser Thr Ser Arg Ser Ser Pro Asn Lys Arg
 1525 1530 1535

 Gly Pro Pro Thr Tyr Asn Glu His Ile Thr Lys Arg Val Ala Ser Ser
 1540 1545 1550

Pro Ala Pro Pro Glu Gly Pro Ser His Pro Arg Glu Pro Ser Thr Pro
1555 1560 1565

His Arg Tyr Arg Asp Arg Glu Gly Arg Thr Glu Leu Arg Arg Asp Lys
1570 1575 1580

Ser Pro Gly Arg Pro Leu Glu Arg Glu Lys Ser Pro Gly Arg Met Leu
1585 1590 1595 1600

Ser Thr Arg Arg Glu Arg Ser Pro Gly Arg Leu Phe Glu Asp Ser Ser
1605 1610 1615

Arg Gly Arg Leu Pro Ala Gly Ala Val Arg Thr Pro Leu Ser Gln Val
1620 1625 1630

Asn Lys Val Trp Asp Gln Ser Ser Val
1635 1640

<210> 41

<211> 1597

<212> PRT

<213> Mus musculus

<400> 41

Met Leu Leu Gly Glu Glu Ala Met Met Glu Gln Glu Met Thr Arg Leu
1 5 10 15

His Arg Arg Val Ser Glu Val Glu Ala Val Leu Ser Gln Lys Glu Val
20 25 30

Glu Leu Lys Ala Ser Glu Thr Gln Arg Ser Leu Leu Glu Gln Asp Leu
35 40 45

Ala Thr Tyr Ile Thr Glu Cys Ser Ser Leu Lys Arg Ser Leu Glu Gln
50 55 60

Ala Arg Met Glu Val Ser Gln Glu Asp Asp Lys Ala Leu Gln Leu Leu
65 70 75 80

His Asp Ile Arg Glu Gln Ser Arg Lys Leu Gln Glu Ile Lys Glu Gln
85 90 95

Glu Tyr Gln Ala Gln Val Glu Glu Met Arg Leu Met Met Asn Gln Leu
100 105 110

Glu Glu Asp Leu Val Ser Ala Arg Arg Ser Asp Leu Tyr Glu Ser
115 120 125

Glu Leu Arg Glu Ser Arg Leu Ala Ala Glu Glu Phe Lys Arg Lys Ala
130 135 140

Asn Glu Cys Gln His Lys Leu Met Lys Ala Lys Asp Gln Gly Lys Pro
145 150 155 160

Glu Val Gly Glu Tyr Ser Lys Leu Glu Lys Ile Asn Ala Glu Gln Gln

165	170	175
Leu Lys Ile Gln Glu Leu Gln Glu Lys Leu Glu Lys Ala Val Lys Ala 180	185	190
Ser Thr Glu Ala Thr Glu Leu Leu Gln Asn Ile Arg Gln Ala Lys Glu 195	200	205
Arg Ala Glu Arg Glu Leu Glu Lys Leu His Asn Arg Glu Asp Ser Ser 210	215	220
Glu Gly Ile Lys Lys Lys Leu Val Glu Ala Glu Glu Arg Arg His Ser 225	230	235
Leu Glu Asn Lys Val Lys Arg Leu Glu Thr Met Glu Arg Arg Glu Asn 245	250	255
Arg Leu Lys Asp Asp Ile Gln Thr Lys Ser Glu Gln Ile Gln Gln Met 260	265	270
Ala Asp Lys Ile Leu Glu Leu Glu Lys His Arg Glu Ala Gln Val 275	280	285
Ser Ala Gln His Leu Glu Val His Leu Lys Gln Lys Glu Gln His Tyr 290	295	300
Glu Glu Lys Ile Lys Val Leu Asp Asn Gln Ile Lys Lys Asp Leu Ala 305	310	315
Asp Lys Glu Ser Leu Glu Asn Met Met Gln Arg His Glu Glu Ala 325	330	335
His Glu Lys Gly Lys Ile Leu Ser Glu Gln Lys Ala Met Ile Asn Ala 340	345	350
Met Asp Ser Lys Ile Arg Ser Leu Glu Gln Arg Ile Val Glu Leu Ser 355	360	365
Glu Ala Asn Lys Leu Ala Ala Asn Ser Ser Leu Phe Thr Gln Arg Asn 370	375	380
Met Lys Ala Gln Glu Glu Met Ile Ser Glu Leu Arg Gln Gln Lys Phe 385	390	395
Tyr Leu Glu Thr Gln Ala Gly Lys Leu Glu Ala Gln Asn Arg Lys Leu 405	410	415
Glu Glu Gln Leu Glu Lys Ile Ser His Gln Asp His Ser Asp Lys Ser 420	425	430
Arg Leu Leu Glu Leu Glu Thr Arg Leu Arg Glu Val Ser Leu Glu His 435	440	445
Glu Glu Gln Lys Leu Glu Leu Lys Arg Gln Leu Thr Glu Leu Gln Leu 450	455	460
Ser Leu Gln Glu Arg Glu Ser Gln Leu Thr Ala Leu Gln Ala Ala Arg		

465	470	475	480
Ala Ala Leu Glu Ser Gln Leu Arg Gln Ala Lys Thr Glu Leu Glu Glu			
485	490	495	
Thr Thr Ala Glu Ala Glu Glu Ile Gln Ala Leu Thr Ala His Arg			
500	505	510	
Asp Glu Ile Gln Arg Lys Phe Asp Ala Leu Arg Asn Ser Cys Thr Val			
515	520	525	
Ile Thr Asp Leu Glu Glu Gln Leu Asn Gln Leu Thr Glu Asp Asn Ala			
530	535	540	
Glu Leu Asn Asn Gln Asn Phe Tyr Leu Ser Lys Gln Leu Asp Glu Ala			
545	550	555	560
Ser Gly Ala Asn Asp Glu Ile Val Gln Leu Arg Ser Glu Val Asp His			
565	570	575	
Leu Arg Arg Glu Ile Thr Glu Arg Glu Met Gln Leu Thr Ser Gln Lys			
580	585	590	
Gln Thr Met Glu Ala Leu Lys Thr Thr Cys Thr Met Leu Glu Glu Gln			
595	600	605	
Val Leu Asp Leu Glu Ala Leu Asn Asp Glu Leu Leu Glu Lys Glu Arg			
610	615	620	
Gln Trp Glu Ala Trp Arg Ser Val Leu Gly Asp Glu Lys Ser Gln Phe			
625	630	635	640
Glu Cys Arg Val Arg Glu Leu Gln Arg Met Leu Asp Thr Glu Lys Gln			
645	650	655	
Ser Arg Ala Arg Ala Asp Gln Arg Ile Thr Glu Ser Arg Gln Val Val			
660	665	670	
Glu Leu Ala Val Lys Glu His Lys Ala Glu Ile Leu Ala Leu Gln Gln			
675	680	685	
Ala Leu Lys Glu Gln Lys Leu Lys Ala Glu Ser Leu Ser Asp Lys Leu			
690	695	700	
Asn Asp Leu Glu Lys Lys His Ala Met Leu Glu Met Asn Ala Arg Ser			
705	710	715	720
Leu Gln Gln Lys Leu Glu Thr Glu Arg Glu Leu Lys Gln Arg Leu Leu			
725	730	735	
Glu Glu Gln Ala Lys Leu Gln Gln Met Asp Leu Gln Lys Asn His			
740	745	750	
Ile Phe Arg Leu Thr Gln Gly Leu Gln Glu Ala Leu Asp Arg Ala Asp			
755	760	765	
Leu Leu Lys Thr Glu Arg Ser Asp Leu Glu Tyr Gln Leu Glu Asn Ile			

770

775

780

Gln Val Leu Tyr Ser His Glu Lys Val Lys Met Glu Gly Thr Ile Ser
785 790 795 800

Gln Gln Thr Lys Leu Ile Asp Phe Leu Gln Ala Lys Met Asp Gln Pro
805 810 815

Ala Lys Lys Lys Lys Val Pro Leu Gln Tyr Asn Glu Leu Lys Leu Ala
820 825 830

Leu Glu Lys Glu Lys Ala Arg Cys Ala Glu Leu Glu Glu Ala Leu Gln
835 840 845

Lys Thr Arg Ile Glu Leu Arg Ser Ala Arg Glu Glu Ala Ala His Arg
850 855 860

Lys Ala Thr Asp His Pro His Pro Ser Thr Pro Ala Thr Ala Arg Gln
865 870 875 880

Gln Ile Ala Met Ser Ala Ile Val Arg Ser Pro Glu His Gln Pro Ser
885 890 895

Ala Met Ser Leu Leu Ala Pro Pro Ser Ser Arg Arg Lys Glu Ser Ser
900 905 910

Thr Pro Glu Glu Phe Ser Arg Arg Leu Lys Glu Arg Met His His Asn
915 920 925

Ile Pro His Arg Phe Asn Val Gly Leu Asn Met Arg Ala Thr Lys Cys
930 935 940

Ala Val Cys Leu Asp Thr Val His Phe Gly Arg Gln Ala Ser Lys Cys
945 950 955 960

Leu Glu Cys Gln Val Met Cys His Pro Lys Cys Ser Thr Cys Leu Pro
965 970 975

Ala Thr Cys Gly Leu Pro Ala Glu Tyr Ala Thr His Phe Thr Glu Ala
980 985 990

Phe Cys Arg Asp Lys Met Asn Ser Pro Gly Leu Gln Ser Lys Glu Pro
995 1000 1005

Gly Ser Ser Leu His Leu Glu Gly Trp Met Lys Val Pro Arg Asn Asn
1010 1015 1020

Lys Arg Gly Gln Gln Gly Trp Asp Arg Lys Tyr Ile Val Leu Glu Gly
1025 1030 1035 1040

Ser Lys Val Leu Ile Tyr Asp Asn Glu Ala Arg Glu Ala Gly Gln Arg
1045 1050 1055

Pro Val Glu Glu Phe Glu Leu Cys Leu Pro Asp Gly Asp Val Ser Ile
1060 1065 1070

His Gly Ala Val Gly Ala Ser Glu Leu Ala Asn Thr Ala Lys Ala Asp

1075	1080	1085
Val Pro Tyr Ile Leu Lys Met Glu Ser His Pro His Thr Thr Cys Trp		
1090	1095	1100
Pro Gly Arg Thr Leu Tyr Leu Leu Ala Pro Ser Phe Pro Asp Lys Gln		
1105	1110	1115
Arg Trp Val Thr Ala Leu Glu Ser Val Val Ala Gly Gly Arg Val Ser		
1125	1130	1135
Arg Glu Lys Ala Glu Ala Asp Ala Lys Leu Leu Gly Asn Ser Leu Leu		
1140	1145	1150
Lys Leu Glu Gly Asp Asp Arg Leu Asp Met Asn Cys Thr Leu Pro Phe		
1155	1160	1165
Ser Asp Gln Val Val Leu Val Gly Thr Glu Glu Gly Leu Tyr Ala Leu		
1170	1175	1180
Asn Val Leu Lys Asn Ser Leu Thr His Ile Pro Gly Ile Gly Ala Val		
1185	1190	1195
Phe Gln Ile Tyr Ile Ile Lys Asp Leu Glu Lys Leu Leu Met Ile Ala		
1205	1210	1215
Gly Glu Glu Arg Ala Leu Cys Leu Val Asp Val Lys Lys Val Lys Gln		
1220	1225	1230
Ser Leu Ala Gln Ser His Leu Pro Ala Gln Pro Asp Val Ser Pro Asn		
1235	1240	1245
Ile Phe Glu Ala Val Lys Gly Cys His Leu Phe Ala Ala Gly Lys Ile		
1250	1255	1260
Glu Asn Ser Leu Cys Ile Cys Ala Ala Met Pro Ser Lys Val Val Ile		
1265	1270	1275
Leu Arg Tyr Asn Asp Asn Leu Ser Lys Tyr Cys Ile Arg Lys Glu Ile		
1285	1290	1295
Glu Thr Ser Glu Pro Cys Ser Cys Ile His Phe Thr Asn Tyr Ser Ile		
1300	1305	1310
Leu Ile Gly Thr Asn Lys Phe Tyr Glu Ile Asp Met Lys Gln Tyr Thr		
1315	1320	1325
Leu Asp Glu Phe Leu Asp Lys Asn Asp His Ser Leu Ala Pro Ala Val		
1330	1335	1340
Phe Ala Ser Ser Ser Asn Ser Phe Pro Val Ser Ile Val Gln Ala Asn		
1345	1350	1355
Ser Ala Gly Gln Arg Glu Glu Tyr Leu Leu Cys Phe His Glu Phe Gly		
1365	1370	1375
Val Phe Val Asp Ser Tyr Gly Arg Arg Ser Arg Thr Asp Asp Leu Lys		

1380	1385	1390
Trp Ser Arg Leu Pro Leu Ala Phe Ala Tyr Arg Glu Pro Tyr Leu Phe		
1395	1400	1405
Val Thr His Phe Asn Ser Leu Glu Val Ile Glu Ile Gln Ala Arg Ser		
1410	1415	1420
Ser Leu Gly Ser Pro Ala Arg Ala Tyr Leu Glu Ile Pro Asn Pro Arg		
1425	1430	1435
Tyr Leu Gly Pro Ala Ile Ser Ser Gly Ala Ile Tyr Leu Ala Ser Ser		
1445	1450	1455
Tyr Gln Asp Lys Leu Arg Val Ile Cys Cys Lys Gly Asn Leu Val Lys		
1460	1465	1470
Glu Ser Gly Thr Glu Gln His Arg Val Pro Ser Thr Ser Arg Ser Ser		
1475	1480	1485
Pro Asn Lys Arg Gly Pro Pro Thr Tyr Asn Glu His Ile Thr Lys Arg		
1490	1495	1500
Val Ala Ser Ser Pro Ala Pro Pro Glu Gly Pro Ser His Pro Arg Glu		
1505	1510	1515
Pro Ser Thr Pro His Arg Tyr Arg Asp Arg Glu Gly Arg Thr Glu Leu		
1525	1530	1535
Arg Arg Asp Lys Ser Pro Gly Arg Pro Leu Glu Arg Glu Lys Ser Pro		
1540	1545	1550
Gly Arg Met Leu Ser Thr Arg Arg Glu Arg Ser Pro Gly Arg Leu Phe		
1555	1560	1565
Glu Asp Ser Ser Arg Gly Arg Leu Pro Ala Gly Ala Val Arg Thr Pro		
1570	1575	1580
Leu Ser Gln Val Asn Lys Val Trp Asp Gln Ser Ser Val		
1585	1590	1595
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Ala Leu Gly Asn Arg Pro Phe Arg Ala Phe Val Val Thr Asp Thr Thr		
20	25	30
Leu Thr His Leu Ala Val His Arg Val Thr Gly Glu Val Phe Val Gly		
35	40	45

Ala Val Asn Arg Val Phe Lys Leu Ala Pro Asn Leu Thr Glu Leu Arg
 50 55 60

Ala His Val Thr Gly Pro Val Glu Asp Asn Ala Arg Cys Tyr Pro Pro
 65 70 75 80

Pro Ser Met Arg Val Cys Ala His Arg Leu Ala Pro Val Asp Asn Ile
 85 90 95

Asn Lys Leu Leu Leu Ile Asp Tyr Ala Ala Arg Arg Leu Val Ala Cys
 100 105 110

Gly Ser Ile Trp Gln Gly Ile Cys Gln Phe Leu Arg Leu Asp Asp Leu
 115 120 125

Phe Lys Leu Gly Glu Pro His His Arg Lys Glu His Tyr Leu Ser Gly
 130 135 140

Ala Gln Glu Pro Asp Ser Met Ala Gly Val Ile Val Glu Gln Gly Gln
 145 150 155 160

Gly Pro Ser Lys Leu Phe Val Gly Thr Ala Val Asp Gly Lys Ser Glu
 165 170 175

Tyr Phe Pro Thr Leu Ser Ser Arg Lys Leu Ile Ser Asp Glu Asp Ser
 180 185 190

Ala Asp Met Phe Ser Leu Val Tyr Gln Asp Glu Phe Val Ser Ser Gln
 195 200 205

Ile Lys Ile Pro Ser Asp Thr Leu Ser Leu Tyr Pro Ala Phe Asp Ile
 210 215 220

Tyr Tyr Ile Tyr Gly Phe Val Ser Ala Ser Phe Val Tyr Phe Leu Thr
 225 230 235 240

Leu Gln Leu Asp Thr Gln Gln Thr Leu Leu Asp Thr Ala Gly Glu Lys
 245 250 255

Phe Phe Thr Ser Lys Ile Val Arg Met Cys Ala Gly Asp Ser Glu Phe
 260 265 270

Tyr Ser Tyr Val Glu Phe Pro Ile Gly Cys Ser Trp Arg Gly Val Glu
 275 280 285

Tyr Arg Leu Val Gln Ser Ala His Leu Ala Lys Pro Gly Leu Leu Leu
 290 295 300

Ala Gln Ala Leu Gly Val Pro Ala Asp Glu Asp Val Leu Phe Thr Ile
 305 310 315 320

Phe Ser Gln Gly Gln Lys Asn Arg Ala Ser Pro Pro Arg Gln Thr Ile
 325 330 335

Leu Cys Leu Phe Thr Leu Ser Asn Ile Asn Ala His Ile Arg Arg Arg
 340 345 350

Ile	Gln	Ser	Cys	Tyr	Arg	Gly	Glu	Gly	Thr	Leu	Ala	Leu	Pro	Trp	Leu
355															365
Leu	Asn	Lys	Glu	Leu	Pro	Cys	Ile	Asn	Thr	Pro	Met	Gln	Ile	Asn	Gly
370															380
Asn	Phe	Cys	Gly	Leu	Val	Leu	Asn	Gln	Pro	Leu	Gly	Gly	Leu	His	Val
385															400
Ile	Glu	Gly	Leu	Pro	Leu	Leu	Ala	Asp	Ser	Thr	Asp	Gly	Met	Ala	Ser
405															415
Val	Ala	Ala	Tyr	Thr	Tyr	Arg	Gln	His	Ser	Val	Val	Phe	Ile	Gly	Thr
420															430
Arg	Ser	Gly	Ser	Leu	Lys	Lys	Val	Arg	Val	Asp	Gly	Phe	Gln	Asp	Ala
435															445
His	Leu	Tyr	Glu	Thr	Val	Pro	Val	Val	Asp	Gly	Ser	Pro	Ile	Leu	Arg
450															460
Asp	Leu	Leu	Phe	Ser	Pro	Asp	His	Arg	His	Ile	Tyr	Leu	Leu	Ser	Glu
465															480
Lys	Gln	Val	Ser	Gln	Leu	Pro	Val	Glu	Thr	Cys	Glu	Gln	Tyr	Gln	Ser
485															495
Cys	Ala	Ala	Cys	Leu	Gly	Ser	Gly	Asp	Pro	His	Cys	Gly	Trp	Cys	Val
500															510
Leu	Arg	His	Arg	Cys	Cys	Arg	Glu	Gly	Ala	Cys	Leu	Gly	Ala	Ser	Ala
515															525
Pro	His	Gly	Phe	Ala	Glu	Glu	Leu	Ser	Lys	Cys	Val	Gln	Val	Arg	Val
530															540
Arg	Pro	Asn	Asn	Val	Ser	Val	Thr	Ser	Pro	Gly	Val	Gln	Leu	Thr	Val
545															560
Thr	Leu	His	Asn	Val	Pro	Asp	Leu	Ser	Ala	Gly	Val	Ser	Cys	Ala	Phe
565															575
Glu	Ala	Ala	Ala	Glu	Asn	Glu	Ala	Val	Leu	Leu	Pro	Ser	Gly	Glu	Leu
580															590
Leu	Cys	Pro	Ser	Pro	Ser	Leu	Gln	Glu	Leu	Arg	Ala	Leu	Thr	Arg	Gly
595															605
His	Gly	Ala	Thr	Arg	Thr	Val	Arg	Leu	Gln	Leu	Leu	Ser	Lys	Glu	Thr
610															620
Gly	Val	Arg	Phe	Ala	Gly	Ala	Asp	Phe	Val	Phe	Tyr	Asn	Cys	Ser	Val
625															640
Leu	Gln	Ser	Cys	Met	Ser	Cys	Val	Gly	Ser	Pro	Tyr	Pro	Cys	His	Trp
645															655

Cys Lys Tyr Arg His Thr Cys Thr Ser Arg Pro His Glu Cys Ser Phe
 660 665 670
 Gln Glu Gly Arg Val His Ser Pro Glu Gly Cys Pro Glu Ile Leu Pro
 675 680 685
 Ser Gly Asp Leu Leu Ile Pro Val Gly Val Met Gln Pro Leu Thr Leu
 690 695 700
 Arg Ala Lys Asn Leu Pro Gln Pro Gln Ser Gly Gln Lys Asn Tyr Glu
 705 710 715 720
 Cys Val Val Arg Val Gln Gly Arg Gln Gln Arg Val Pro Ala Val Arg
 725 730 735
 Phe Asn Ser Ser Ser Val Gln Cys Gln Asn Ala Ser Tyr Ser Tyr Glu
 740 745 750
 Gly Asp Glu His Gly Asp Thr Glu Leu Asp Phe Ser Val Val Trp Asp
 755 760 765
 Gly Asp Phe Pro Ile Asp Lys Pro Pro Ser Phe Arg Ala Leu Leu Tyr
 770 775 780
 Lys Cys Trp Ala Gln Arg Pro Ser Cys Gly Leu Cys Leu Lys Ala Asp
 785 790 795 800
 Pro Arg Phe Asn Cys Gly Trp Cys Ile Ser Glu His Arg Cys Gln Leu
 805 810 815
 Arg Thr His Cys Pro Ala Pro Lys Thr Asn Trp Met His Leu Ser Gln
 820 825 830
 Lys Gly Thr Arg Cys Ser His Pro Arg Ile Thr Gln Ile His Pro Leu
 835 840 845
 Val Gly Pro Lys Glu Gly Thr Arg Val Thr Ile Val Gly Asp Asn
 850 855 860
 Leu Gly Leu Leu Ser Arg Glu Val Gly Leu Arg Val Ala Gly Val Arg
 865 870 875 880
 Cys Asn Ser Ile Pro Ala Glu Tyr Ile Ser Ala Glu Arg Ile Val Cys
 885 890 895
 Glu Met Glu Glu Ser Leu Val Pro Ser Pro Pro Pro Gly Pro Val Glu
 900 905 910
 Leu Cys Val Gly Asp Cys Ser Ala Asp Phe Arg Thr Gln Ser Glu Gln
 915 920 925
 Val Tyr Ser Phe Val Thr Pro Thr Phe Asp Gln Val Ser Pro Ser Arg
 930 935 940
 Gly Pro Ala Ser Gly Gly Thr Arg Leu Thr Ile Ser Gly Ser Ser Leu
 945 950 955 960

Asp Ala Gly Ser Arg Val Thr Val Thr Val Arg Asp Ser Glu Cys Gln
 965 970 975

 Phe Val Arg Arg Asp Ala Lys Ala Ile Val Cys Ile Ser Pro Leu Ser
 980 985 990

 Thr Leu Gly Pro Ser Gln Ala Pro Ile Thr Leu Ala Ile Asp Arg Ala
 995 1000 1005

 Asn Ile Ser Ser Pro Gly Leu Ile Tyr Thr Tyr Thr Gln Asp Pro Thr
 1010 1015 1020

 Val Thr Arg Leu Glu Pro Thr Trp Ser Ile Ile Asn Gly Ser Thr Ala
 1025 1030 1035 1040

 Ile Thr Val Ser Gly Thr His Leu Leu Thr Val Gln Glu Pro Arg Val
 1045 1050 1055

 Arg Ala Lys Tyr Arg Gly Ile Glu Thr Thr Asn Thr Cys Gln Val Ile
 1060 1065 1070

 Asn Asp Thr Ala Met Leu Cys Lys Ala Pro Gly Ile Phe Leu Gly Arg
 1075 1080 1085

 Pro Gln Pro Arg Ala Gln Gly Glu His Pro Asp Glu Phe Gly Phe Leu
 1090 1095 1100

 Leu Asp His Val Gln Thr Ala Arg Ser Leu Asn Arg Ser Ser Phe Thr
 1105 1110 1115 1120

 Tyr Tyr Pro Asp Pro Ser Phe Glu Pro Leu Gly Pro Ser Gly Val Leu
 1125 1130 1135

 Asp Val Lys Pro Gly Ser His Val Val Leu Lys Gly Lys Asn Leu Ile
 1140 1145 1150

 Pro Ala Ala Ala Gly Ser Ser Arg Leu Asn Tyr Thr Val Leu Ile Gly
 1155 1160 1165

 Gly Gln Pro Cys Ser Leu Thr Val Ser Asp Thr Gln Leu Leu Cys Asp
 1170 1175 1180

 Ser Pro Ser Gln Thr Gly Arg Gln Pro Val Met Val Leu Val Gly Gly
 1185 1190 1195 1200

 Leu Glu Phe Trp Leu Gly Thr Leu His Ile Ser Ala Glu Arg Ala Leu
 1205 1210 1215

 Thr Leu Pro Ala Met Met Gly Leu Ala Ala Gly Gly Leu Leu Leu
 1220 1225 1230

 Leu Ala Ile Thr Ala Val Leu Val Ala Tyr Lys Arg Lys Thr Gln Asp
 1235 1240 1245

 Ala Asp Arg Thr Leu Lys Arg Leu Gln Leu Gln Met Asp Asn Leu Glu
 1250 1255 1260

Ser Arg Val Ala Leu Glu Cys Lys Glu Ala Phe Ala Glu Leu Gln Thr
 1265 1270 1275 1280

 Asp Ile Asn Glu Leu Thr Asn His Met Asp Glu Val Gln Ile Pro Phe
 1285 1290 1295

 Leu Asp Tyr Arg Thr Tyr Ala Val Arg Val Leu Phe Pro Gly Ile Glu
 1300 1305 1310

 Ala His Pro Val Leu Lys Glu Leu Asp Thr Pro Pro Asn Val Glu Lys
 1315 1320 1325

 Ala Leu Arg Leu Phe Gly Gln Leu Leu His Ser Arg Ala Phe Val Leu
 1330 1335 1340

 Thr Phe Ile His Thr Leu Glu Ala Gln Ser Ser Phe Ser Met Arg Asp
 1345 1350 1355 1360

 Arg Gly Thr Val Ala Ser Leu Thr Met Val Ala Leu Gln Ser Arg Leu
 1365 1370 1375

 Asp Tyr Ala Thr Gly Leu Leu Lys Gln Leu Leu Ala Asp Leu Ile Glu
 1380 1385 1390

 Lys Asn Leu Glu Ser Lys Asn His Pro Lys Leu Leu Leu Arg Arg Thr
 1395 1400 1405

 Glu Ser Val Ala Glu Lys Met Leu Thr Asn Trp Phe Thr Phe Leu Leu
 1410 1415 1420

 His Lys Phe Leu Lys Glu Cys Ala Gly Glu Pro Leu Phe Leu Leu Tyr
 1425 1430 1435 1440

 Cys Ala Ile Lys Gln Gln Met Glu Lys Gly Pro Ile Asp Ala Ile Thr
 1445 1450 1455

 Gly Glu Ala Arg Tyr Ser Leu Ser Glu Asp Lys Leu Ile Arg Gln Gln
 1460 1465 1470

 Ile Asp Tyr Lys Thr Leu Thr Leu His Cys Val Cys Pro Glu Asn Glu
 1475 1480 1485

 Gly Ser Ala Gln Val Pro Val Lys Val Leu Asn Cys Asp Ser Ile Thr
 1490 1495 1500

 Gln Ala Lys Asp Lys Leu Leu Asp Thr Val Tyr Lys Gly Ile Pro Tyr
 1505 1510 1515 1520

 Ser Gln Arg Pro Lys Ala Glu Asp Met Asp Leu Glu Trp Arg Gln Gly
 1525 1530 1535

 Arg Met Thr Arg Ile Ile Leu Gln Asp Glu Asp Val Thr Thr Lys Ile
 1540 1545 1550

 Glu Cys Asp Trp Lys Arg Leu Asn Ser Leu Ala His Tyr Gln Val Thr
 1555 1560 1565

Asp Gly Ser Leu Val Ala Leu Val Pro Lys Gln Val Ser Ala Tyr Asn
1570 1575 1580

Met Ala Asn Ser Phe Thr Phe Thr Arg Ser Leu Ser Arg Tyr Glu Ser
1585 1590 1595 1600

Leu Leu Arg Thr Ala Ser Ser Pro Asp Ser Leu Arg Ser Arg Ala Pro
1605 1610 1615

Met Ile Thr Pro Asp Gln Glu Thr Gly Thr Lys Leu Trp His Leu Val
1620 1625 1630

Lys Asn His Asp His Ala Asp His Arg Glu Gly Asp Arg Gly Ser Lys
1635 1640 1645

Met Val Ser Glu Ile Tyr Leu Thr Arg Leu Leu Ala Thr Lys Gly Thr
1650 1655 1660

Leu Gln Lys Phe Val Asp Asp Leu Phe Glu Thr Val Phe Ser Thr Ala
1665 1670 1675 1680

His Arg Gly Ser Ala Leu Pro Leu Ala Ile Lys Tyr Met Phe Asp Phe
1685 1690 1695

Leu Asp Glu Gln Ala Asp Gln Arg Gln Ile Ser Asp Pro Asp Val Arg
1700 1705 1710

His Thr Trp Lys Ser Asn Cys Leu Pro Leu Arg Phe Trp Val Asn Val
1715 1720 1725

Ile Lys Asn Pro Gln Phe Val Phe Asp Ile His Lys Asn Ser Ile Thr
1730 1735 1740

Asp Ala Cys Leu Ser Val Val Ala Gln Thr Phe Met Asp Ser Cys Ser
1745 1750 1755 1760

Thr Ser Glu His Arg Leu Gly Lys Asp Ser Pro Ser Asn Lys Leu Leu
1765 1770 1775

Tyr Ala Lys Asp Ile Pro Asn Tyr Lys Ser Trp Val Glu Arg Tyr Tyr
1780 1785 1790

Arg Asp Ile Ala Lys Met Ala Ser Ile Ser Asp Gln Asp Met Asp Ala
1795 1800 1805

Tyr Leu Val Glu Gln Ser Arg Leu His Ala Ser Asp Phe Ser Val Leu
1810 1815 1820

Ser Ala Leu Asn Glu Leu Tyr Phe Tyr Val Thr Lys Tyr Arg Gln Glu
1825 1830 1835 1840

Ile Leu Thr Ala Leu Asp Arg Asp Ala Ser Cys Arg Lys His Lys Leu
1845 1850 1855

Arg Gln Lys Leu Glu Gln Ile Ile Ser Leu Val Ser Ser Asp Ser
1860 1865 1870

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<212> PRT
<213> Homo sapiens

<400> 43
Gly Phe Gly Arg Leu Pro Asp Ser Glu Leu Arg Ala Gly Arg Gly Ala
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Ser Arg Arg Pro Gln Gln Pro Ala Ala Ala Glu Val Asp Arg Ala Gly
20 25 30

Thr Glu Gly Gln Thr Asp Val Ala Glu Leu Glu Ser Cys Glu Gly Gln
35 40 45

Pro Gly Lys Val Glu Gln Met Ser Thr His Arg Ser Arg Leu Leu Thr
50 55 60

Ala Ala Pro Leu Ser Met Glu Gln Arg Arg Pro Trp Pro Arg Ala Leu
65 70 75 80

Glu Val Asp Ser Arg Ser Val Val Leu Leu Ser Val Val Trp Val Leu
85 90 95

Leu Ala Pro Pro Ala Ala Gly Met Pro Gln Phe Ser Thr Phe His Ser
100 105 110

Glu Asn Arg Asp Trp Thr Phe Asn His Leu Thr Val His Gln Gly Thr
115 120 125

Gly Ala Val Tyr Val Gly Ala Ile Asn Arg Val Tyr Lys Leu Thr Gly
130 135 140

Asn Leu Thr Ile Gln Val Ala His Lys Thr Gly Pro Glu Glu Asp Asn
145 150 155 160

Lys Ser Cys Tyr Pro Pro Leu Ile Val Gln Pro Cys Ser Glu Val Leu
165 170 175

Thr Leu Thr Asn Asn Val Asn Lys Leu Leu Ile Ile Asp Tyr Ser Glu
180 185 190

Asn Arg Leu Leu Ala Cys Gly Ser Leu Tyr Gln Gly Val Cys Lys Leu
195 200 205

Leu Arg Leu Asp Asp Leu Phe Ile Leu Val Glu Pro Ser His Lys Lys
210 215 220

Glu His Tyr Leu Ser Ser Val Asn Lys Thr Gly Thr Met Tyr Gly Val
225 230 235 240

Ile Val Arg Ser Glu Gly Glu Asp Gly Lys Leu Phe Ile Gly Thr Ala
245 250 255

Val Asp Gly Lys Gln Asp Tyr Phe Pro Thr Leu Ser Ser Arg Lys Leu
260 265 270

Pro Arg Asp Pro Glu Ser Ser Ala Met Leu Asp Tyr Glu Leu His Ser
 275 280 285

 Asp Phe Val Ser Ser Leu Ile Lys Ile Pro Ser Asp Thr Leu Ala Leu
 290 295 300

 Val Ser His Phe Asp Ile Phe Tyr Ile Tyr Gly Phe Ala Ser Gly Gly
 305 310 315 320

 Phe Val Tyr Phe Leu Thr Val Gln Pro Glu Thr Pro Glu Gly Val Ala
 325 330 335

 Ile Asn Ser Ala Gly Asp Leu Phe Tyr Thr Ser Arg Ile Val Arg Leu
 340 345 350

 Cys Lys Asp Asp Pro Lys Phe His Ser Tyr Val Ser Leu Pro Phe Gly
 355 360 365

 Cys Thr Arg Ala Gly Val Glu Tyr Arg Leu Leu Gln Ala Ala Tyr Leu
 370 375 380

 Ala Lys Pro Gly Asp Ser Leu Ala Gln Ala Phe Asn Ile Thr Ser Gln
 385 390 395 400

 Asp Asp Val Leu Phe Ala Ile Phe Ser Lys Gly Gln Lys Gln Tyr His
 405 410 415

 His Pro Pro Asp Asp Ser Ala Leu Cys Ala Phe Pro Ile Arg Ala Ile
 420 425 430

 Asn Leu Gln Ile Lys Gly Arg Leu Gln Ser Cys Tyr Gln Gly Glu Gly
 435 440 445

 Asn Leu Glu Leu Asn Trp Leu Leu Gly Lys Asp Val Gln Cys Thr Lys
 450 455 460

 Ala Pro Val Pro Ile Asp Asp Asn Phe Cys Gly Leu Asp Ile Asn Gln
 465 470 475 480

 Pro Leu Gly Gly Ser Thr Pro Val Glu Gly Leu Thr Leu Tyr Thr Thr
 485 490 495

 Ser Arg Asp Arg Met Thr Ser Val Ala Ser Tyr Val Tyr Asn Gly Tyr
 500 505 510

 Ser Val Val Phe Val Gly Thr Lys Ser Gly Lys Leu Lys Lys Ile Arg
 515 520 525

 Ala Asp Gly Pro Pro His Gly Gly Val Gln Tyr Glu Met Val Ser Val
 530 535 540

 Leu Lys Asp Gly Ser Pro Ile Leu Arg Asp Met Ala Phe Ser Ile Asp
 545 550 555 560

 Gln Arg Tyr Leu Tyr Val Met Ser Glu Arg Gln Val Thr Arg Val Pro
 565 570 575

Val	Glu	Ser	Cys	Glu	Gln	Tyr	Thr	Thr	Cys	Gly	Glu	Cys	Leu	Ser	Ser
580							585							590	
Gly	Asp	Pro	His	Cys	Gly	Trp	Cys	Ala	Leu	His	Asn	Met	Cys	Ser	Arg
595							600							605	
Arg	Asp	Lys	Cys	Gln	Gln	Ala	Trp	Glu	Pro	Asn	Arg	Phe	Ala	Ala	Ser
610							615							620	
Ile	Ser	Gln	Cys	Val	Ser	Leu	Ala	Val	His	Pro	Ser	Ser	Ile	Ser	Val
625							630				635			640	
Ser	Glu	His	Ser	Arg	Leu	Leu	Ser	Leu	Val	Val	Ser	Asp	Ala	Pro	Asp
645							650							655	
Leu	Ser	Ala	Gly	Ile	Ala	Cys	Ala	Phe	Gly	Asn	Leu	Thr	Glu	Val	Glu
660							665							670	
Gly	Gln	Val	Ser	Gly	Ser	Gln	Val	Ile	Cys	Ile	Ser	Pro	Gly	Pro	Lys
675							680							685	
Asp	Val	Pro	Val	Ile	Pro	Leu	Asp	Gln	Asp	Trp	Phe	Gly	Leu	Glu	Leu
690							695							700	
Gln	Leu	Arg	Ser	Lys	Glu	Thr	Gly	Lys	Ile	Phe	Val	Ser	Thr	Glu	Phe
705							710				715			720	
Lys	Phe	Tyr	Asn	Cys	Ser	Ala	His	Gln	Leu	Cys	Leu	Ser	Cys	Val	Asn
725							730							735	
Ser	Ala	Phe	Arg	Cys	His	Trp	Cys	Lys	Tyr	Arg	Asn	Leu	Cys	Thr	His
740							745							750	
Asp	Pro	Thr	Thr	Cys	Ser	Phe	Gln	Glu	Gly	Arg	Ile	Asn	Ile	Ser	Glu
755							760							765	
Asp	Cys	Pro	Gln	Leu	Val	Pro	Thr	Glu	Glu	Ile	Leu	Ile	Pro	Val	Gly
770							775							780	
Glu	Val	Lys	Pro	Ile	Thr	Leu	Lys	Ala	Arg	Asn	Leu	Pro	Gln	Pro	Gln
785							790				795			800	
Ser	Gly	Gln	Arg	Gly	Tyr	Glu	Cys	Val	Leu	Asn	Ile	Gln	Gly	Ala	Ile
805							810							815	
His	Arg	Val	Pro	Ala	Leu	Arg	Phe	Asn	Ser	Ser	Ser	Val	Gln	Cys	Gln
820							825							830	
Asn	Ser	Ser	Tyr	Gln	Tyr	Asp	Gly	Met	Asp	Ile	Ser	Asn	Leu	Ala	Val
835							840							845	
Asp	Phe	Ala	Val	Val	Trp	Asn	Gly	Asn	Phe	Ile	Ile	Asp	Asn	Pro	Gln
850							855							860	
Asp	Leu	Lys	Val	His	Leu	Tyr	Lys	Cys	Ala	Ala	Gln	Arg	Glu	Ser	Cys
865							870				875			880	

Gly Leu Cys Leu Lys Ala Asp Arg Lys Phe Glu Cys Gly Trp Cys Ser
885 890 895

Gly Glu Arg Arg Cys Thr Leu His Gln His Cys Thr Ser Pro Ser Ser
900 905 910

Pro Trp Leu Asp Trp Ser Ser His Asn Val Lys Cys Ser Asn Pro Gln
915 920 925

Ile Thr Glu Ile Leu Thr Val Ser Gly Pro Pro Glu Gly Gly Thr Arg
930 935 940

Val Thr Ile His Gly Val Asn Leu Gly Leu Asp Phe Ser Glu Ile Ala
945 950 955 960

His His Val Gln Val Ala Gly Val Pro Cys Thr Pro Leu Pro Gly Glu
965 970 975

Tyr Ile Ile Ala Glu Gln Ile Val Cys Glu Met Gly His Ala Leu Val
980 985 990

Gly Thr Thr Ser Gly Pro Val Arg Leu Cys Ile Gly Glu Cys Lys Pro
995 1000 1005

Glu Phe Met Thr Lys Ser His Gln Gln Tyr Thr Phe Val Asn Pro Ser
1010 1015 1020

Val Leu Ser Leu Asn Pro Ile Arg Gly Pro Glu Ser Gly Gly Thr Met
1025 1030 1035 1040

Val Thr Ile Thr Gly His Tyr Leu Gly Ala Gly Ser Ser Val Ala Val
1045 1050 1055

Tyr Leu Gly Asn Gln Thr Cys Glu Phe Tyr Gly Arg Ser Met Ser Glu
1060 1065 1070

Ile Val Cys Val Ser Pro Pro Ser Ser Asn Gly Leu Gly Pro Val Pro
1075 1080 1085

Val Ser Val Ser Val Asp Arg Ala His Val Asp Ser Asn Leu Gln Phe
1090 1095 1100

Glu Tyr Ile Asp Asp Pro Arg Val Gln Arg Ile Glu Pro Glu Trp Ser
1105 1110 1115 1120

Ile Ala Ser Gly His Thr Pro Leu Thr Ile Thr Gly Phe Asn Leu Asp
1125 1130 1135

Val Ile Gln Glu Pro Arg Ile Arg Val Lys Phe Asn Gly Lys Glu Ser
1140 1145 1150

Val Asn Val Cys Lys Val Val Asn Thr Thr Leu Thr Cys Leu Ala
1155 1160 1165

Pro Ser Leu Thr Thr Asp Tyr Arg Pro Gly Leu Asp Thr Val Glu Arg
1170 1175 1180

Pro Asp Glu Phe Gly Phe Val Phe Asn Asn Val Gln Ser Leu Leu Ile
 1185 1190 1195 1200
 Tyr Asn Asp Thr Lys Phe Ile Tyr Tyr Pro Asn Pro Thr Phe Glu Leu
 1205 1210 1215
 Leu Ser Pro Thr Gly Val Leu Asp Gln Lys Pro Gly Ser Pro Ile Ile
 1220 1225 1230
 Leu Lys Gly Lys Asn Leu Cys Pro Pro Ala Ser Gly Gly Ala Lys Leu
 1235 1240 1245
 Asn Tyr Thr Val Leu Ile Gly Glu Thr Pro Cys Ala Val Thr Val Ser
 1250 1255 1260
 Glu Thr Gln Leu Leu Cys Glu Pro Pro Asn Leu Thr Gly Gln His Lys
 1265 1270 1275 1280
 Val Met Val His Val Gly Gly Met Val Phe Ser Pro Gly Ser Val Ser
 1285 1290 1295
 Val Ile Ser Asp Ser Leu Leu Thr Leu Pro Ala Ile Val Ser Ile Ala
 1300 1305 1310
 Ala Gly Gly Ser Leu Leu Leu Ile Val Ile Ile Val Leu Ile Ala
 1315 1320 1325
 Tyr Lys Arg Lys Ser Arg Glu Asn Asp Leu Thr Leu Lys Arg Leu Gln
 1330 1335 1340
 Met Gln Met Asp Asn Leu Glu Ser Arg Val Ala Leu Glu Cys Lys Glu
 1345 1350 1355 1360
 Ala Phe Ala Glu Leu Gln Thr Asp Ile Asn Glu Leu Thr Ser Asp Leu
 1365 1370 1375
 Asp Arg Ser Gly Ile Pro Tyr Leu Asp Tyr Arg Thr Tyr Ala Met Arg
 1380 1385 1390
 Val Leu Phe Pro Gly Ile Glu Asp His Pro Val Leu Arg Glu Leu Glu
 1395 1400 1405
 Val Gln Gly Asn Gly Gln Gln His Val Glu Lys Ala Leu Lys Leu Phe
 1410 1415 1420
 Ala Gln Leu Ile Asn Asn Lys Val Phe Leu Leu Thr Phe Ile Arg Thr
 1425 1430 1435 1440
 Leu Glu Leu Gln Arg Ser Phe Ser Met Arg Asp Arg Gly Asn Val Ala
 1445 1450 1455
 Ser Leu Ile Met Thr Gly Leu Gln Gly Arg Leu Glu Tyr Ala Thr Asp
 1460 1465 1470
 Val Leu Lys Gln Leu Leu Ser Asp Leu Ile Asp Lys Asn Leu Glu Asn
 1475 1480 1485

Lys Asn His Pro Lys Leu Leu Leu Arg Arg Thr Glu Ser Val Ala Glu
1490 1495 1500

Lys Met Leu Thr Asn Trp Phe Ala Phe Leu Leu His Lys Phe Leu Lys
1505 1510 1515 1520

Glu Cys Ala Gly Glu Pro Leu Phe Met Leu Tyr Cys Ala Ile Lys Gln
1525 1530 1535

Gln Met Glu Lys Gly Pro Ile Asp Ala Ile Thr Gly Glu Ala Arg Tyr
1540 1545 1550

Ser Leu Ser Glu Asp Lys Leu Ile Arg Gln Gln Ile Glu Tyr Lys Thr
1555 1560 1565

Leu Ile Leu Asn Cys Val Asn Pro Asp Asn Glu Asn Ser Pro Glu Ile
1570 1575 1580

Pro Val Lys Val Leu Asn Cys Asp Thr Ile Thr Gln Val Lys Glu Lys
1585 1590 1595 1600

Ile Leu Asp Ala Val Tyr Lys Asn Val Pro Tyr Ser Gln Arg Pro Arg
1605 1610 1615

Ala Val Asp Met Asp Leu Glu Trp Arg Gln Gly Arg Ile Ala Arg Val
1620 1625 1630

Val Leu Gln Asp Glu Asp Ile Thr Thr Lys Ile Glu Gly Asp Trp Lys
1635 1640 1645

Arg Leu Asn Thr Leu Met His Tyr Gln Val Ser Asp Arg Ser Val Val
1650 1655 1660

Ala Leu Val Pro Lys Gln Thr Ser Ser Tyr Asn Ile Pro Ala Ser Ala
1665 1670 1675 1680

Ser Ile Ser Arg Thr Ser Ile Ser Arg Tyr Asp Ser Ser Phe Arg Tyr
1685 1690 1695

Thr Gly Ser Pro Asp Ser Leu Arg Ser Arg Ala Pro Met Ile Thr Pro
1700 1705 1710

Asp Leu Glu Ser Gly Val Lys Val Trp His Leu Val Lys Asn His Asp
1715 1720 1725

His Gly Asp Gln Lys Glu Gly Asp Arg Gly Ser Lys Met Val Ser Glu
1730 1735 1740

Ile Tyr Leu Thr Arg Leu Leu Ala Thr Lys Gly Thr Leu Gln Lys Phe
1745 1750 1755 1760

Val Asp Asp Leu Phe Glu Thr Leu Phe Ser Thr Val His Arg Gly Ser
1765 1770 1775

Ala Leu Pro Leu Ala Ile Lys Tyr Met Phe Asp Phe Leu Asp Glu Gln
1780 1785 1790

Ala Asp Arg His Ser Ile His Asp Thr Asp Val Arg His Thr Trp Lys
1795 1800 1805

Ser Asn Cys Leu Pro Leu Arg Phe Trp Val Asn Val Ile Lys Asn Pro
1810 1815 1820

Gln Phe Val Phe Asp Ile His Lys Gly Ser Ile Thr Asp Ala Cys Leu
1825 1830 1835 1840

Ser Val Val Ala Gln Thr Phe Met Asp Ser Cys Ser Thr Ser Glu His
1845 1850 1855

Arg Leu Gly Lys Asp Ser Pro Ser Asn Lys Leu Leu Tyr Ala Lys Asp
1860 1865 1870

Ile Pro Ser Tyr Lys Ser Trp Val Glu Arg Tyr Tyr Ala Asp Ile Ala
1875 1880 1885

Lys Leu Pro Ala Ile Ser Asp Gln Asp Met Asn Ala Tyr Leu Ala Glu
1890 1895 1900

Gln Ser Arg Leu His Ala Val Glu Phe Asn Met Leu Ser Ala Leu Asn
1905 1910 1915 1920

Glu Ile Tyr Ser Tyr Val Ser Lys Tyr Ser Glu Glu Leu Ile Gly Ala
1925 1930 1935

Leu Glu Gln Asp Glu Gln Ala Arg Arg Gln Arg Leu Ala Tyr Lys Val
1940 1945 1950

Glu Gln Leu Ile Asn Ala Met Ser Ile Glu Ser
1955 1960

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<211> 1905
<212> PRT
<213> Xenopus laevis

<400> 44
Met Leu Leu His Ala Glu Arg Pro Leu Pro Phe His Leu Trp Thr Phe
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Leu Val Leu Leu Gly Ser Trp Ile Ala Thr Gly Asp Gly Ser Pro Lys
20 25 30

Asp Phe Arg Thr Phe Thr Gly Ser Asp Trp Ser Leu Thr His Leu Val
35 40 45

Val His Asn Lys Thr Gly Glu Val Tyr Val Gly Ala Ile Asn Arg Ile
50 55 60

Tyr Lys Leu Ser Asn Asn Leu Thr Leu Leu Arg Thr His Val Thr Gly
65 70 75 80

Pro Val Glu Asp Asn Glu Lys Cys Tyr Pro Pro Pro Ser Val Gln Ser

85	90	95
Cys Pro His Gly Leu Ile Thr Thr Asn Asn Val Asn Lys Leu Leu Leu		
100	105	110
Ile Asp Tyr Ser Asp Asn Arg Leu Ile Ala Cys Gly Ser Ala Ser Gln		
115	120	125
Gly Ile Cys Gln Phe Leu Arg Leu Asp Asp Leu Phe Lys Leu Gly Glu		
130	135	140
Pro His His Arg Lys Glu His Tyr Leu Ser Ser Val Asn Glu Ser Gly		
145	150	155
160		
Thr Met Ser Gly Val Ile Ile Glu Val Pro Asn Gly Gln Asn Lys Leu		
165	170	175
Phe Val Gly Thr Pro Ile Asp Gly Lys Ser Glu Tyr Phe Pro Thr Leu		
180	185	190
Ser Ser Arg Lys Leu Leu Gly Asn Glu Glu Asn Ala Glu Met Phe Gly		
195	200	205
Phe Val Tyr Gln Asp Glu Phe Val Ser Ser Gln Leu Lys Ile Pro Ser		
210	215	220
Asp Thr Leu Ser Lys Phe Pro Thr Phe Asp Ile Tyr Tyr Val Tyr Ser		
225	230	235
240		
Phe Ser Ser Glu Gln Phe Val Tyr Tyr Leu Thr Leu Gln Leu Asp Thr		
245	250	255
Gln Leu Thr Ser Pro Asp Ser Thr Gly Glu Gln Phe Phe Thr Ser Lys		
260	265	270
Ile Val Arg Leu Cys Val Asp Asp Pro Lys Phe Tyr Ser Tyr Val Glu		
275	280	285
Phe Pro Ile Gly Cys Met Lys Asp Gly Val Glu Tyr Arg Leu Ile Gln		
290	295	300
Asp Ala Tyr Leu Ser Lys Pro Gly Lys Arg Leu Ala Lys Glu Leu Gly		
305	310	315
320		
Ile Ser Glu Arg Glu Asp Ile Leu Phe Thr Val Phe Ser Gln Gly Gln		
325	330	335
Lys Asn Arg Ile Lys Pro Pro Lys Glu Ser Val Leu Cys Leu Phe Thr		
340	345	350
Leu Lys Lys Ile Lys Asp Lys Ile Lys Glu Arg Ile Gln Ser Cys Tyr		
355	360	365
Arg Gly Asp Gly Lys Leu Ser Leu Pro Trp Leu Leu Asn Lys Glu Leu		
370	375	380
Gly Cys Ile Asn Ser Pro Leu Gln Ile Asp Asp Asn Phe Cys Gly Gln		

385	390	395	400
Asp Phe Asn Gln Pro Leu Gly Gly Thr Val Thr Ile Glu Gly Thr Pro			
405	410	415	
Leu Phe Leu Asp Lys Glu Asp Gly Met Thr Ser Val Ala Ala Tyr Asp			
420	425	430	
Tyr Arg Gly His Thr Val Val Phe Ala Gly Thr Arg Ser Gly Arg Val			
435	440	445	
Lys Lys Ile Leu Val Asp Leu Ser Ala Ser Ser Ser His Leu Val Gln			
450	455	460	
Gln Tyr Glu Asn Val Val Val His Glu Gly Asn Ala Ile Leu Arg Asp			
465	470	475	480
Leu Val Leu Ser Pro Asp Arg Gln Tyr Ile Tyr Ala Met Thr Glu Lys			
485	490	495	
Gln Val Thr Arg Val Pro Val Glu Ser Cys Glu Gln Tyr Glu Ser Cys			
500	505	510	
Asp Thr Cys Leu Gly Ser Arg Asp Pro His Cys Gly Trp Cys Val Leu			
515	520	525	
His Asn Met Cys Ser Arg Lys Asp Lys Cys Glu Arg Ala Asp Glu Leu			
530	535	540	
His Arg Phe Thr Ser Asp Gln Arg Gln Cys Val Gln Leu Thr Val His			
545	550	555	560
Pro Lys Asn Ile Ser Val Thr Val Ser Glu Val Pro Met Val Leu Gln			
565	570	575	
Ala Trp Asn Val Pro Asp Leu Ser Ala Gly Val Asn Cys Ser Phe Glu			
580	585	590	
Asp Phe Thr Glu Met Glu Gly Arg Ile Leu Asp Gly Lys Ile Tyr Cys			
595	600	605	
Thr Ser Pro Ser Ala Lys Glu Val Ile Pro Ile Thr Arg Gly His Gly			
610	615	620	
Asp Lys Arg Val Val Lys Leu Tyr Leu Lys Ser Lys Glu Thr Gly Lys			
625	630	635	640
Lys Phe Ala Ser Val Asp Phe Val Phe Tyr Asn Cys Ser Val His Gln			
645	650	655	
Ser Cys Leu Ser Cys Val Asn Gly Ser Phe Pro Cys His Trp Cys Lys			
660	665	670	
Tyr Arg His Val Cys Thr His Asn Ala Ala Asp Cys Ser Phe Gln Glu			
675	680	685	
Gly Arg Val Asn Met Ser Glu Asp Cys Pro Gln Ile Leu Pro Ser Ser			

690

695

700

Gln Ile Tyr Ile Pro Val Gly Val Val Lys Pro Ile Thr Leu Thr Ala
 705 710 715 720

Lys Asn Leu Pro Gln Pro Gln Ser Gly Gln Arg Asn Tyr Glu Cys Ile
 725 730 735

Phe His Ile Pro Gly Ser Val Thr Arg Val Thr Ala Leu Arg Phe Asn
 740 745 750

Ser Thr Ser Ile Gln Cys Gln Asn Thr Ser Tyr Asn Tyr Glu Gly Asn
 755 760 765

Asp Ile Ser Asp Leu Pro Val Asn Leu Ser Val Val Trp Asn Gly His
 770 775 780

Phe Val Ile Asp Asn Pro Gln Asn Ile Gln Ala His Leu Tyr Lys Cys
 785 790 795 800

Ser Ala Leu Arg Glu Ser Cys Gly Leu Cys Leu Lys Ser Asp Arg Arg
 805 810 815

Phe Glu Cys Gly Trp Cys Val Ser Glu Lys Lys Cys Thr Leu Arg Gln
 820 825 830

Asn Cys Pro Thr Leu Glu Asn Pro Trp Met His Ala Ser Thr Ala Asn
 835 840 845

Ser Arg Cys Thr Asp Pro Lys Ile Thr Lys Leu Phe Pro Glu Thr Gly
 850 855 860

Pro Arg Gln Gly Gly Thr Arg Leu Thr Ile Thr Gly Glu Asn Leu Gly
 865 870 875 880

Leu Arg Phe Glu Asp Ile Arg Phe Gly Val Arg Val Gly His Val Met
 885 890 895

Cys Val Pro Val Glu Ser Glu Tyr Ile Ser Ala Glu Gln Ile Val Cys
 900 905 910

Glu Ile Asn Asp Ala Gly Arg Thr Arg Val His Glu Ala Gln Val Glu
 915 920 925

Val Cys Val Lys Asp Cys Ser Gln Asp Tyr Arg Ala Ile Ser Pro Lys
 930 935 940

Ser Phe Thr Phe Val Leu Pro Ser Phe Asn Arg Val Thr Pro Ser Arg
 945 950 955 960

Gly Pro Leu Ser Gly Gly Thr Trp Ile Ser Ile Glu Gly Asn Tyr Leu
 965 970 975

Asn Ala Gly Ser Asp Val Ser Val Ala Ile Gly Gly Arg Pro Cys Met
 980 985 990

Phe Ser Trp Arg Thr Ala Lys Glu Ile Arg Cys Lys Thr Pro Gln Gly

995	1000	1005
Pro Ser Thr Gly Lys Ala Glu Ile Gln Ile Leu Ile Asn Arg Ala Thr		
1010	1015	1020
Met Asn Asn Ser Glu Val His Tyr Asn Tyr Thr Glu Asp Pro Thr Val		
1025	1030	1035
1040		
Gln Lys Ile Glu Pro Glu Trp Ser Ile Ala Ser Gly Gly Thr Pro Leu		
1045	1050	1055
Ile Val Thr Gly Met Asn Leu Ala Thr Ile Lys Glu Pro Lys Ile Arg		
1060	1065	1070
Ala Lys Tyr Gly Asp Val Glu Lys Glu Asn Asn Cys Thr Leu Tyr Asn		
1075	1080	1085
Asp Thr Thr Met Val Cys Leu Ala Pro Ser Val Asp Asn Pro Leu Arg		
1090	1095	1100
Ser Pro Pro Glu Asn Gly Asp Arg Pro Asp Glu Ile Gly Phe Ile Met		
1105	1110	1115
1120		
Asp Asn Val His Ala Leu Leu Ile Val Asn Thr Thr Ser Phe Leu Tyr		
1125	1130	1135
Tyr Pro Asp Pro Val Phe Glu Pro Leu Thr Ala Ser Gly Asn Leu Glu		
1140	1145	1150
Leu Lys Pro Ser Ser Pro Leu Ile Ile Lys Gly Arg Asn Leu Ile Pro		
1155	1160	1165
Ala Ala Pro Gly Asn Phe Arg Leu Asn Tyr Thr Val Leu Ile Gly Asp		
1170	1175	1180
Thr Pro Cys Ala Leu Thr Val Ser Glu Thr Gln Leu Leu Cys Glu Ser		
1185	1190	1195
1200		
Pro Asn Leu Thr Gly Gln His Lys Val Thr Ile Lys Ala Gly Gly Phe		
1205	1210	1215
Glu Tyr Ser Pro Gly Thr Leu Gln Ile Tyr Ser Asp Ser Leu Leu Thr		
1220	1225	1230
Leu Pro Ala Ile Ile Gly Ile Gly Gly Gly Leu Leu Leu Leu		
1235	1240	1245
Ile Ile Ile Ile Val Leu Ile Ala Tyr Lys Arg Lys Ser Arg Asp Ala		
1250	1255	1260
Asp Arg Thr Leu Lys Arg Leu Gln Leu Gln Met Asp Asn Leu Glu Ser		
1265	1270	1275
1280		
Arg Val Ala Leu Glu Cys Lys Glu Ala Phe Ala Glu Leu Gln Thr Asp		
1285	1290	1295
Ile His Glu Leu Thr Asn Asp Leu Asp Gly Ala Gly Ile Pro Phe Leu		

	1300	1305	1310
Glu Tyr Arg Thr Tyr Ala Met Arg Val Leu Phe Pro Gly Ile Glu Asp			
1315	1320	1325	
His Pro Val Leu Lys Glu Met Glu Val Gln Ala Asn Val Glu Lys Ser			
1330	1335	1340	
Leu Thr Leu Phe Gly Gln Leu Leu Thr Lys Lys His Phe Leu Leu Thr			
1345	1350	1355	1360
Phe Ile Arg Thr Leu Glu Ala Gln Arg Ser Phe Ser Met Arg Asp Arg			
1365	1370	1375	
Gly Asn Val Ala Ser Leu Ile Met Thr Ala Leu Gln Gly Glu Met Glu			
1380	1385	1390	
Tyr Ala Thr Gly Val Leu Lys Gln Leu Leu Ser Asp Leu Ile Glu Lys			
1395	1400	1405	
Asn Leu Glu Ser Lys Asn His Pro Lys Leu Leu Leu Arg Arg Thr Glu			
1410	1415	1420	
Ser Val Ala Glu Lys Met Leu Thr Asn Trp Phe Thr Phe Leu Leu Tyr			
1425	1430	1435	1440
Lys Phe Leu Lys Glu Cys Ala Gly Glu Pro Leu Phe Met Leu His Cys			
1445	1450	1455	
Ala Ile Lys Gln Gln Met Glu Lys Gly Pro Ile Asp Ala Ile Thr Gly			
1460	1465	1470	
Glu Ala Arg Tyr Ser Leu Ser Glu Asp Lys Leu Ile Arg Gln Gln Ile			
1475	1480	1485	
Asp Tyr Lys Thr Leu Asn Pro Cys Ala Asp Asp Val Gly Leu Ser Asp			
1490	1495	1500	
Glu Ser Cys Cys Arg Ser Pro Gln Thr Leu Asn Cys Val Asn Pro Glu			
1505	1510	1515	1520
Asn Glu Asn Ala Pro Glu Ile Pro Val Lys Val Leu Asn Cys Asp Thr			
1525	1530	1535	
Ile Thr Gln Val Lys Glu Lys Leu Leu Asp Ala Val Tyr Lys Gly Val			
1540	1545	1550	
Pro Tyr Ser Gln Arg Pro Lys Ala Gly Asp Met Asp Leu Glu Trp Arg			
1555	1560	1565	
Gln Gly Arg Met Ala Arg Ile Ile Leu Gln Asp Glu Asp Val Thr Thr			
1570	1575	1580	
Lys Ile Asp Asn Asp Trp Lys Arg Leu Asn Thr Leu Ala His Tyr Gln			
1585	1590	1595	1600
Val Thr Asp Gly Ser Ser Val Ala Leu Val Pro Lys Gln Asn Ser Ala			

1605	1610	1615	
Tyr Asn Ile Ser Asn Ser Ser Thr Phe Thr Lys Ser Leu Ser Arg Tyr			
1620	1625	1630	
Glu Ser Met Leu Arg Thr Ala Ser Ser Pro Asp Ser Leu Arg Ser Arg			
1635	1640	1645	
Thr Pro Met Ile Thr Pro Asp Leu Glu Ser Gly Thr Lys Leu Trp His			
1650	1655	1660	
Leu Val Lys Asn His Asp His Leu Asp Gln Arg Glu Gly Asp Arg Gly			
1665	1670	1675	1680
Ser Lys Met Val Ser Glu Ile Tyr Leu Thr Arg Leu Leu Ala Thr Lys			
1685	1690	1695	
Gly Thr Leu Gln Lys Phe Val Asp Asp Leu Phe Glu Thr Ile Phe Ser			
1700	1705	1710	
Thr Ala His Arg Gly Ser Ala Leu Pro Leu Ala Ile Lys Tyr Met Phe			
1715	1720	1725	
Asp Phe Leu Asp Glu Gln Ala Asp Lys His Gln Ile Thr Asp Tyr Asp			
1730	1735	1740	
Val Arg His Thr Trp Lys Ser Asn Cys Leu Pro Leu Arg Phe Trp Val			
1745	1750	1755	1760
Asn Val Ile Lys Asn Pro Gln Phe Val Phe Asp Ile His Lys Asn Ser			
1765	1770	1775	
Ile Thr Asp Ala Cys Leu Ser Val Val Ala Gln Thr Phe Met Asp Ser			
1780	1785	1790	
Cys Ser Thr Ser Glu His Lys Leu Gly Lys Asp Ser Pro Ser Asn Lys			
1795	1800	1805	
Leu Leu Tyr Ala Lys Asp Ile Pro Asn Tyr Lys Ser Trp Val Glu Arg			
1810	1815	1820	
Tyr Tyr Ala Asp Ile Ala Lys Met Pro Val Ile Ser Asp Gln Asp Met			
1825	1830	1835	1840
Ser Ala Tyr Leu Ala Glu Gln Ser Arg Leu His Leu Ser Gln Phe Asn			
1845	1850	1855	
Ser Met Ser Ala Leu His Glu Ile Tyr Ser Tyr Ile Thr Lys Tyr Arg			
1860	1865	1870	
Asp Glu Ile Leu Thr Ala Leu Glu Lys Asp Glu Gln Ala Arg Arg Gln			
1875	1880	1885	
Arg Leu Arg Ser Lys Leu Glu Gln Val Ile Asp Thr Met Ala Gln Ser			
1890	1895	1900	
Ser			

<210> 45
<211> 813
<212> PRT
<213> Homo sapiens

<400> 45
Gly Thr Arg Val Val Cys Lys Val Val Asn Thr Thr Thr Leu Thr Cys
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Leu Ala Pro Ser Leu Thr Thr Asp Tyr Arg Pro Gly Leu Asp Thr Val
20 25 30
Glu Arg Pro Asp Glu Phe Gly Phe Val Phe Asn Asn Val Gln Ser Leu
35 40 45
Leu Ile Tyr Asn Asp Thr Lys Phe Ile Tyr Tyr Pro Asn Pro Thr Phe
50 55 60
Glu Leu Leu Ser Pro Thr Gly Val Leu Asp Gln Lys Pro Gly Ser Pro
65 70 75 80
Ile Ile Leu Lys Gly Lys Asn Leu Cys Pro Pro Ala Ser Gly Gly Ala
85 90 95
Lys Leu Asn Tyr Thr Val Leu Ile Gly Glu Thr Pro Cys Ala Val Thr
100 105 110
Val Ser Glu Thr Gln Leu Leu Cys Glu Pro Pro Asn Leu Thr Gly Gln
115 120 125
His Lys Val Met Val His Val Gly Gly Met Val Phe Ser Pro Gly Ser
130 135 140
Val Ser Val Ile Ser Asp Ser Leu Leu Thr Leu Pro Ala Ile Val Ser
145 150 155 160
Ile Ala Ala Gly Gly Ser Leu Leu Leu Ile Ile Val Ile Val Leu
165 170 175
Ile Ala Tyr Lys Arg Lys Ser Arg Glu Asn Asp Leu Thr Leu Lys Arg
180 185 190
Leu Gln Met Gln Met Asp Asn Leu Glu Ser Arg Val Ala Leu Glu Cys
195 200 205
Lys Glu Ala Phe Ala Glu Leu Gln Thr Asp Ile Asn Glu Leu Thr Ser
210 215 220
Asp Leu Asp Arg Ser Gly Ile Pro Tyr Leu Asp Tyr Arg Thr Tyr Ala
225 230 235 240
Met Arg Val Leu Phe Pro Gly Ile Glu Asp His Pro Val Leu Arg Glu
245 250 255

Leu Glu Val Gln Gly Asn Gly Gln His Val Glu Lys Ala Leu Lys
260 265 270

Leu Phe Ala Gln Leu Ile Asn Asn Lys Val Phe Leu Leu Thr Phe Ile
275 280 285

Arg Thr Leu Glu Leu Gln Arg Ser Phe Ser Met Arg Asp Arg Gly Asn
290 295 300

Val Ala Ser Leu Ile Met Thr Gly Leu Gln Gly Arg Leu Glu Tyr Ala
305 310 315 320

Thr Asp Val Leu Lys Gln Leu Leu Ser Asp Leu Ile Asp Lys Asn Leu
325 330 335

Glu Asn Lys Asn His Pro Lys Leu Leu Arg Arg Thr Glu Ser Val
340 345 350

Ala Glu Lys Met Leu Thr Asn Trp Phe Ala Phe Leu Leu His Lys Phe
355 360 365

Leu Lys Glu Cys Ala Gly Glu Pro Leu Phe Met Leu Tyr Cys Ala Ile
370 375 380

Lys Gln Gln Met Glu Lys Gly Pro Ile Asp Ala Ile Thr Gly Glu Ala
385 390 395 400

Arg Tyr Ser Leu Ser Glu Asp Lys Leu Ile Arg Gln Gln Ile Glu Tyr
405 410 415

Lys Thr Leu Ile Leu Asn Cys Val Asn Pro Asp Asn Glu Asn Ser Pro
420 425 430

Glu Ile Pro Val Lys Val Leu Asn Cys Asp Thr Ile Thr Gln Val Lys
435 440 445

Glu Lys Ile Leu Asp Ala Val Tyr Lys Asn Val Pro Tyr Ser Gln Arg
450 455 460

Pro Arg Ala Val Asp Met Asp Leu Glu Trp Arg Gln Gly Arg Ile Ala
465 470 475 480

Arg Val Val Leu Gln Asp Glu Asp Ile Thr Thr Lys Ile Glu Gly Asp
485 490 495

Trp Lys Arg Leu Asn Thr Leu Met His Tyr Gln Val Ser Asp Arg Ser
500 505 510

Val Val Ala Leu Val Pro Lys Gln Thr Ser Ser Tyr Asn Ile Pro Ala
515 520 525

Ser Ala Ser Ile Ser Arg Thr Ser Ile Ser Arg Tyr Asp Ser Ser Phe
530 535 540

Arg Tyr Thr Gly Ser Pro Asp Ser Leu Arg Ser Arg Ala Pro Met Ile
545 550 555 560

Thr Pro Asp Leu Glu Ser Gly Val Lys Val Trp His Leu Val Lys Asn
 565 570 575

 His Asp His Gly Asp Gln Lys Glu Gly Asp Arg Gly Ser Lys Met Val
 580 585 590

 Ser Glu Ile Tyr Leu Thr Arg Leu Leu Ala Thr Lys Gly Thr Leu Gln
 595 600 605

 Lys Phe Val Asp Asp Leu Phe Glu Thr Leu Leu Ser Thr Val His Arg
 610 615 620

 Gly Ser Ala Leu Pro Leu Ala Ile Lys Tyr Met Phe Asp Phe Leu Asp
 625 630 635 640

 Glu Gln Ala Asp Arg His Ser Ile His Asp Thr Asp Val Arg His Thr
 645 650 655

 Trp Lys Ser Asn Cys Leu Pro Leu Arg Phe Trp Val Asn Val Ile Lys
 660 665 670

 Asn Pro Gln Phe Val Phe Asp Ile His Lys Gly Ser Ile Thr Asp Ala
 675 680 685

 Cys Leu Ser Val Val Ala Gln Thr Phe Met Asp Ser Cys Ser Thr Ser
 690 695 700

 Glu His Arg Leu Gly Lys Asp Ser Pro Ser Asn Lys Leu Leu Tyr Ala
 705 710 715 720

 Lys Asp Ile Pro Ser Tyr Lys Ser Trp Val Glu Arg Tyr Tyr Ala Asp
 725 730 735

 Ile Ala Lys Leu Pro Ala Ile Ser Asp Gln Asp Met Asn Ala Tyr Leu
 740 745 750

 Ala Glu Gln Ser Arg Leu His Ala Val Glu Phe Asn Met Leu Ser Ala
 755 760 765

 Leu Asn Glu Ile Tyr Ser Tyr Val Ser Lys Tyr Ser Glu Glu Leu Ile
 770 775 780

 Gly Ala Leu Glu Gln Asp Glu Gln Ala Arg Arg Gln Arg Leu Ala Tyr
 785 790 795 800

 Lys Val Glu Gln Leu Ile Asn Ala Met Ser Ile Glu Ser
 805 810

<210> 46
 <211> 593
 <212> PRT
 <213> Homo sapiens

<400> 46
 Val Glu Leu Thr Val Val Trp Asn Gly His Phe Asn Ile Asp Asn Pro
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Ala Gln Asn Lys Val His Leu Tyr Lys Cys Gly Ala Met Arg Glu Ser
20 25 30

Cys Gly Leu Cys Leu Lys Ala Asp Pro Asp Phe Ala Cys Gly Trp Cys
35 40 45

Gln Gly Pro Gly Gln Cys Thr Leu Arg Gln His Cys Pro Ala Gln Glu
50 55 60

Ser Gln Trp Leu Glu Leu Ser Gly Ala Lys Ser Lys Cys Thr Asn Pro
65 70 75 80

Arg Ile Thr Glu Ile Ile Pro Val Thr Gly Pro Arg Glu Gly Gly Thr
85 90 95

Lys Val Thr Ile Arg Gly Glu Asn Leu Gly Leu Glu Phe Arg Asp Ile
100 105 110

Ala Ser His Val Lys Val Ala Gly Val Glu Cys Ser Pro Leu Val Asp
115 120 125

Gly Tyr Ile Pro Ala Glu Gln Ile Val Cys Glu Met Gly Glu Ala Lys
130 135 140

Pro Ser Gln His Ala Gly Phe Val Glu Ile Cys Val Ala Val Cys Arg
145 150 155 160

Pro Glu Phe Met Ala Arg Ser Ser Gln Leu Tyr Tyr Phe Met Thr Leu
165 170 175

Thr Leu Ser Asp Leu Lys Pro Ser Arg Gly Pro Met Ser Gly Gly Thr
180 185 190

Gln Val Thr Ile Thr Gly Thr Asn Leu Asn Ala Gly Ser Asn Val Val
195 200 205

Val Met Phe Gly Lys Gln Pro Cys Leu Phe His Arg Arg Ser Pro Ser
210 215 220 240

Tyr Ile Val Cys Asn Thr Thr Ser Ser Asp Glu Val Leu Glu Met Lys
225 230 235 240

Val Ser Val Gln Val Asp Arg Ala Lys Ile His Gln Asp Leu Val Phe
245 250 255

Gln Tyr Val Glu Asp Pro Thr Ile Val Arg Ile Glu Pro Glu Trp Ser
260 265 270

Ile Val Ser Gly Asn Thr Pro Ile Ala Val Trp Gly Thr His Leu Asp
275 280 285

Leu Ile Gln Asn Pro Gln Ile Arg Ala Lys His Gly Gly Lys Glu His
290 295 300

Ile Asn Ile Cys Glu Val Leu Asn Ala Thr Glu Met Thr Cys Gln Ala
305 310 315 320

Pro Ala Leu Ala Leu Gly Pro Asp His Gln Ser Asp Leu Thr Glu Arg
325 330 335

Pro Glu Glu Phe Gly Phe Ile Leu Asp Asn Val Gln Ser Leu Leu Ile
340 345 350

Leu Asn Lys Thr Asn Phe Thr Tyr Tyr Pro Asn Pro Val Phe Glu Ala
355 360 365

Phe Gly Pro Ser Gly Ile Leu Glu Leu Lys Pro Gly Thr Pro Ile Ile
370 375 380

Leu Lys Gly Lys Asn Leu Ile Pro Pro Val Ala Gly Gly Asn Val Lys
385 390 395 400

Leu Asn Tyr Thr Val Leu Val Gly Glu Lys Pro Cys Thr Val Thr Val
405 410 415

Ser Asp Val Gln Leu Leu Cys Glu Ser Pro Asn Leu Ile Gly Arg His
420 425 430

Lys Val Met Ala Arg Val Gly Gly Met Glu Tyr Ser Pro Gly Met Val
435 440 445

Tyr Ile Ala Pro Asp Ser Pro Leu Ser Leu Pro Ala Ile Val Ser Ile
450 455 460

Ala Val Ala Gly Gly Leu Leu Ile Ile Phe Ile Val Ala Val Leu Ile
465 470 475 480

Ala Tyr Lys Arg Lys Ser Arg Glu Ser Asp Leu Thr Leu Lys Arg Leu
485 490 495

Gln Met Gln Met Asp Asn Leu Glu Ser Arg Val Ala Leu Glu Cys Lys
500 505 510

Glu Gly Thr Glu Trp Pro His Ala Gly Gly His Val Cys Val Arg Val
515 520 525

Cys Ile Cys Val Cys Met His Ile Cys Val Cys Val Cys Ile Cys Phe
530 535 540

Ile Tyr Lys Gln Ala Gly Trp Ala Ala Val Gly Ser Ala Gly Gly Trp
545 550 555 560

Arg Cys Val Cys Leu Cys Glu Cys Val Cys Val His Val Cys Val Cys
565 570 575

Thr Ser Val Cys Ile Tyr Val Ser Tyr Thr Ser Lys Gln Ala Gly Gln
580 585 590

Gln

<211> 477
<212> PRT
<213> Homo sapiens

<400> 47
Met Leu Pro Pro Gly Ser Asn Gly Thr Ala Tyr Pro Gly Gln Phe Ala
1 5 10 15

Leu Tyr Gln Gln Leu Ala Gln Gly Asn Ala Val Gly Gly Ser Ala Gly
20 25 30

Ala Pro Pro Leu Gly Pro Ser Gln Val Val Thr Ala Cys Leu Leu Thr
35 40 45

Leu Leu Ile Ile Trp Thr Leu Leu Gly Asn Val Leu Val Cys Ala Ala
50 55 60

Ile Val Arg Ser Arg His Leu Arg Ala Asn Met Thr Asn Val Phe Ile
65 70 75 80

Val Ser Leu Ala Val Ser Asp Leu Phe Val Ala Leu Leu Val Met Pro
85 90 95

Trp Lys Ala Val Ala Glu Val Ala Gly Tyr Trp Pro Phe Gly Ala Phe
100 105 110

Cys Asp Val Trp Val Ala Phe Asp Ile Met Cys Ser Thr Ala Ser Ile
115 120 125

Leu Asn Leu Cys Val Ile Ser Val Asp Arg Tyr Trp Ala Ile Ser Arg
130 135 140

Pro Phe Arg Tyr Lys Arg Lys Met Thr Gln Arg Met Ala Leu Val Met
145 150 155 160

Val Gly Leu Ala Trp Thr Leu Ser Ile Leu Ile Ser Phe Ile Pro Val
165 170 175

Gln Leu Asn Trp His Arg Asp Gln Ala Ala Ser Trp Gly Gly Leu Asp
180 185 190

Leu Pro Asn Asn Leu Ala Asn Trp Thr Pro Trp Glu Glu Asp Phe Trp
195 200 205

Glu Pro Asp Val Asn Ala Glu Asn Cys Asp Ser Ser Leu Asn Arg Thr
210 215 220

Tyr Ala Ile Ser Ser Ser Leu Ile Ser Phe Tyr Ile Pro Val Ala Ile
225 230 235 240

Met Ile Val Thr Tyr Thr Arg Ile Tyr Arg Ile Ala Gln Val Gln Ile
245 250 255

Arg Arg Ile Ser Ser Leu Glu Arg Ala Ala Glu His Ala Gln Ser Cys
260 265 270

Arg Ser Ser Ala Ala Cys Ala Pro Asp Thr Ser Leu Arg Ala Ser Ile

275

280

285

Lys Lys Glu Thr Lys Val Leu Lys Thr Leu Ser Val Ile Met Gly Val
 290 295 300

Phe Val Cys Cys Trp Leu Pro Phe Phe Ile Leu Asn Cys Met Val Pro
 305 310 315 320

Phe Cys Ser Gly His Pro Glu Gly Pro Pro Ala Gly Phe Pro Cys Val
 325 330 335

Ser Glu Thr Thr Phe Asp Val Phe Val Trp Phe Gly Trp Ala Asn Ser
 340 345 350

Ser Leu Asn Pro Val Ile Tyr Ala Phe Asn Ala Asp Phe Gln Lys Val
 355 360 365

Phe Ala Gln Leu Leu Gly Cys Ser His Phe Cys Ser Arg Thr Pro Val
 370 375 380

Glu Thr Val Asn Ile Ser Asn Glu Leu Ile Ser Tyr Asn Gln Asp Ile
 385 390 395 400

Val Phe His Lys Glu Ile Ala Ala Tyr Ile His Met Met Pro Asn
 405 410 415

Ala Val Thr Pro Gly Asn Arg Glu Val Asp Asn Asp Glu Glu Gly
 420 425 430

Pro Phe Asp Arg Met Phe Gln Ile Tyr Gln Thr Ser Pro Asp Gly Asp
 435 440 445

Pro Val Ala Glu Ser Val Trp Glu Leu Asp Cys Glu Gly Glu Ile Ser
 450 455 460

Leu Asp Lys Ile Thr Pro Phe Thr Pro Asn Gly Phe His
 465 470 475

<210> 48

<211> 475

<212> PRT

<213> Homo sapiens

<400> 48

Met Leu Pro Pro Gly Arg Asn Arg Thr Ala Gln Pro Ala Arg Leu Gly
 1 5 10 15

Leu Gln Arg Gln Leu Ala Gln Val Asp Ala Pro Ala Gly Ser Ala Thr
 20 25 30

Pro Leu Gly Pro Ala Gln Val Val Thr Ala Gly Leu Leu Thr Leu Leu
 35 40 45

Ile Val Trp Thr Leu Leu Gly Asn Val Leu Val Cys Ala Ala Ile Val
 50 55 60

Arg Ser Arg His Leu Arg Ala Lys Met Thr Asn Ile Phe Ile Val Ser
 65 70 75 80
 Leu Ala Val Ser Asp Leu Phe Val Ala Leu Leu Val Met Pro Trp Lys
 85 90 95
 Ala Val Ala Glu Val Ala Gly Tyr Trp Pro Phe Gly Thr Phe Cys Asp
 100 105 110
 Ile Trp Val Ala Phe Asp Ile Met Cys Ser Thr Ala Ser Ile Leu Asn
 115 120 125
 Leu Cys Ile Ile Ser Val Asp Arg Tyr Trp Ala Ile Ser Arg Pro Phe
 130 135 140
 Arg Tyr Glu Arg Lys Met Thr Gln Arg Val Ala Leu Val Met Val Gly
 145 150 155 160
 Leu Ala Trp Thr Leu Ser Ile Leu Ile Ser Phe Ile Pro Val Gln Leu
 165 170 175
 Asn Trp His Arg Asp Lys Ala Gly Ser Gln Gly Gln Glu Gly Leu Leu
 180 185 190
 Ser Asn Gly Thr Pro Trp Glu Glu Gly Trp Glu Leu Glu Gly Arg Thr
 195 200 205
 Glu Asn Cys Asp Ser Ser Leu Asn Arg Thr Tyr Ala Ile Ser Ser Ser
 210 215 220
 Leu Ile Ser Phe Tyr Ile Pro Val Ala Ile Met Ile Val Thr Tyr Thr
 225 230 235 240
 Arg Ile Tyr Arg Ile Ala Gln Val Gln Ile Arg Arg Ile Ser Ser Leu
 245 250 255
 Glu Arg Ala Ala Glu His Ala Gln Ser Cys Arg Ser Arg Gly Ala Tyr
 260 265 270
 Glu Pro Asp Pro Ser Leu Arg Ala Ser Ile Lys Lys Glu Thr Lys Val
 275 280 285
 Phe Lys Thr Leu Ser Met Ile Met Gly Val Phe Val Cys Cys Trp Leu
 290 295 300
 Pro Phe Phe Ile Leu Asn Cys Met Val Pro Phe Cys Ser Ser Gly Asp
 305 310 315 320
 Ala Glu Gly Pro Lys Thr Gly Phe Pro Cys Val Ser Glu Thr Thr Phe
 325 330 335
 Asp Ile Phe Val Trp Phe Gly Trp Ala Asn Ser Ser Leu Asn Pro Ile
 340 345 350
 Ile Tyr Ala Phe Asn Ala Asp Phe Arg Lys Val Phe Ala Gln Leu Leu
 355 360 365

Gly Cys Ser His Phe Cys Phe Arg Thr Pro Val Gln Thr Val Asn Ile
 370 375 380
 Ser Asn Glu Leu Ile Ser Tyr Asn Gln Asp Thr Val Phe His Lys Glu
 385 390 395 400
 Ile Ala Thr Ala Tyr Val His Met Ile Pro Asn Ala Val Ser Ser Gly
 405 410 415
 Asp Arg Glu Val Gly Glu Glu Glu Gly Pro Phe Asp His Met
 420 425 430
 Ser Gln Ile Ser Pro Thr Thr Pro Asp Gly Asp Leu Ala Ala Glu Ser
 435 440 445
 Val Trp Glu Leu Asp Cys Glu Glu Glu Val Ser Leu Gly Lys Ile Ser
 450 455 460
 Pro Leu Thr Pro Asn Cys Phe Asp Lys Thr Ala
 465 470 475

 <210> 49
 <211> 457
 <212> PRT
 <213> Homo sapiens

 <400> 49
 Met Tyr Gln Pro Phe Gln His Leu Asp Ser Asp Gln Val Ala Ser Trp
 1 5 10 15
 Gln Ser Pro Glu Met Leu Met Asn Lys Ser Val Ser Arg Glu Ser Gln
 20 25 30
 Arg Arg Lys Glu Leu Val Ala Gly Gln Ile Val Thr Gly Ser Leu Leu
 35 40 45
 Leu Leu Leu Ile Phe Trp Thr Leu Phe Gly Asn Ile Leu Val Cys Thr
 50 55 60
 Ala Val Met Arg Phe Arg His Leu Arg Ser Arg Val Thr Asn Ile Phe
 65 70 75 80
 Ile Val Ser Leu Ala Val Ser Asp Leu Leu Val Ala Leu Leu Val Met
 85 90 95
 Pro Trp Lys Ala Val Ala Glu Val Ala Gly His Trp Pro Phe Gly Ala
 100 105 110
 Phe Cys Asp Ile Trp Val Ala Phe Asp Ile Met Cys Ser Thr Ala Ser
 115 120 125
 Ile Leu Asn Leu Cys Val Ile Ser Val Asp Arg Tyr Trp Ala Ile Ser
 130 135 140
 Ser Pro Phe Arg Tyr Glu Arg Lys Met Thr Gln Arg Val Ala Leu Leu
 145 150 155 160

Met Ile Ser Thr Ala Trp Ala Leu Ser Val Leu Ile Ser Phe Ile Pro
 165 170 175

 Val Gln Leu Ser Trp His Lys Ser Glu Thr Glu Asp His Leu Leu Ser
 180 185 190

 Asn His Ser Thr Gly Asn Cys Asp Ser Ser Leu Asn Arg Thr Tyr Ala
 195 200 205

 Ile Ser Ser Ser Leu Ile Ser Phe Tyr Ile Pro Val Ala Ile Met Ile
 210 215 220

 Val Thr Tyr Thr Arg Ile Tyr Arg Ile Ala Gln Ile Gln Ile Lys Arg
 225 230 235 240

 Ile Ser Thr Leu Glu Arg Ala Ala Glu His Ala Gln Ser Cys Arg Ser
 245 250 255

 Asn Arg Val Asp Ser Cys Ser Arg His His Gln Thr Ser Leu Arg Thr
 260 265 270

 Ser Ile Lys Lys Glu Thr Lys Val Leu Lys Thr Leu Ser Ile Ile Met
 275 280 285

 Gly Val Phe Val Cys Cys Trp Leu Pro Phe Phe Ile Leu Asn Cys Met
 290 295 300

 Val Pro Phe Cys Asp Arg Ser Pro Gly His Pro Gln Ala Gly Leu Pro
 305 310 315 320

 Cys Val Ser Glu Thr Thr Phe Asp Ile Phe Val Trp Phe Gly Trp Ala
 325 330 335

 Asn Ser Ser Leu Asn Pro Ile Ile Tyr Ala Phe Asn Ala Asp Phe Arg
 340 345 350

 Lys Val Phe Ser Ser Leu Leu Gly Cys Gly His Trp Cys Ser Thr Thr
 355 360 365

 Pro Val Glu Thr Val Asn Ile Ser Asn Glu Leu Ile Ser Tyr Asn Gln
 370 375 380

 Asp Thr Leu Phe His Lys Asp Ile Val Thr Ala Tyr Val Asn Met Ile
 385 390 395 400

 Pro Asn Val Val Asp Cys Ile Asp Asp Asn Glu Asp Ala Phe Asp His
 405 410 415

 Met Ser Gln Ile Ser Gln Thr Ser Ala Asn Asn Glu Leu Ala Thr Asp
 420 425 430

 Ser Met Cys Glu Leu Asp Ser Glu Val Asp Ile Ser Leu His Lys Ile
 435 440 445

 Thr Pro Ser Met Ser Asn Gly Ile His
 450 455

<210> 50
 <211> 486
 <212> PRT
 <213> Gallus gallus

<400> 50
 Met Leu Arg Gly Gly Arg Ser Pro Leu Pro Pro Pro Ala Gly Pro Pro
 1 5 10 15

Gly Gly Ala Arg Gly Gln Ala Gly Ala Ala Gln Val Ala Ala
 20 25 30

Gly Ser Leu Leu Ala Leu Leu Ile Leu Trp Thr Leu Phe Gly Asn Val
 35 40 45

Leu Val Cys Ala Ala Ile Val Arg Tyr Arg His Leu Arg Ser Lys Val
 50 55 60

Thr Asn Ile Phe Ile Val Ser Leu Ala Val Ser Asp Leu Leu Val Ala
 65 70 75 80

Val Leu Val Met Pro Trp Lys Ala Val Ala Glu Val Ala Gly Tyr Trp
 85 90 95

Pro Phe Gly Ala Phe Gln Asn Val Trp Val Ala Phe Asp Ile Met Cys
 100 105 110

Ser Thr Ala Ser Ile Leu Asn Leu Cys Val Ile Ser Val Asp Arg Tyr
 115 120 125

Trp Ala Ile Ser Ser Pro Phe Arg Tyr Glu Arg Lys Met Thr Gln Arg
 130 135 140

Leu Ala Leu Val Met Ile Gly Val Ala Trp Ala Leu Ser Val Leu Ile
 145 150 155 160

Ser Phe Ile Pro Val Gln Leu Asn Trp His Arg Gly Gly Asp Ala Ala
 165 170 175

Thr Ala Ala Ala Ala Gly Asp Ile Glu Asp Gly Phe Asp Thr Gly Trp
 180 185 190

Glu Ala Ala Gly Ala Phe Thr Thr Trp Ala Glu Asp Met Ser Thr Thr
 195 200 205

Trp Val Ala Leu Ala Ala Met Thr Pro Ser Glu Gly Thr Ser Gly Ser
 210 215 220

Asn Asn Thr Val Pro Gly Pro Ser Glu Ser Cys Asp Ser Ser Leu Asn
 225 230 235 240

Arg Thr Tyr Ala Ile Ser Ser Ser Leu Ile Ser Phe Tyr Ile Pro Val
 245 250 255

Ala Ile Met Ile Val Thr Tyr Thr Arg Ile Tyr Arg Ile Ala Gln Val

260

265

270

Gln Ile Arg Arg Ile Ser Ser Leu Glu Arg Ala Ala Glu His Ala Gln
275 280 285

Ser Cys Arg Cys Asn His Val Asp Cys His His His Thr Ser Leu Lys
290 295 300

Ser Ser Ile Arg Lys Glu Thr Lys Val Leu Lys Thr Leu Ser Ile Ile
305 310 315 320

Met Gly Val Phe Val Cys Cys Trp Leu Pro Phe Phe Ile Leu Asn Cys
325 330 335

Met Val Pro Phe Cys Glu Ser Pro Pro Ser Asp Pro Arg Ala Gly Leu
340 345 350

Pro Cys Val Ser Glu Thr Thr Phe Asn Ile Phe Val Trp Phe Gly Trp
355 360 365

Ala Asn Ser Ser Leu Asn Pro Ile Ile Tyr Ala Phe Asn Ala Asp Phe
370 375 380

Arg Lys Val Phe Ser Asn Leu Leu Gly Cys Gly Gln Phe Cys Ser Ser
385 390 395 400

Thr Pro Val Glu Thr Val Asn Ile Ser Asn Glu Leu Ile Ser Tyr His
405 410 415

Gln Asp Thr Phe His Lys Glu Ile Val Thr Ala Tyr Val Asn Met Ile
420 425 430

Pro Asn Val Val Asp Cys Glu Glu Asn Arg Glu Asp Pro Phe Asp Arg
435 440 445

Met Ser Gln Ile Ser Pro Asp Pro Glu Val Ala Thr Asp Ser Val Cys
450 455 460

Glu Leu Asp Cys Glu Gly Glu Ile Ser Leu Gly Lys Ile Thr Pro Phe
465 470 475 480

Thr Pro Asn Gly Leu His
485

<210> 51

<211> 458

<212> PRT

<213> Anguilla anguilla

<400> 51

Met Gly Ser Pro Ala Lys Tyr Leu Ser Val His Glu Thr Gln Ser Val
1 5 10 15

Pro Phe Phe Ile Gly Glu Ile Met Trp Asn Thr Ser Glu Ser Ala Glu
20 25 30

Lys Thr Asp Gly Lys Lys Glu Leu Ile Val Arg Thr Val Thr Gly Cys
 35 40 45

 Leu Leu Ser Leu Leu Ile Leu Trp Thr Leu Leu Gly Asn Ile Leu Val
 50 55 60

 Cys Ser Ala Val Leu Lys Phe Arg His Leu Arg Thr Lys Val Thr Asn
 65 70 75 80

 Ile Phe Ile Val Ser Leu Ala Val Ser Asp Leu Phe Val Ala Val Leu
 85 90 95

 Val Met Pro Trp Lys Ala Val Ala Glu Val Ala Gly Tyr Trp Pro Phe
 100 105 110

 Gly Pro Phe Cys Asn Ile Trp Val Ala Phe Asp Ile Met Cys Ser Thr
 115 120 125

 Ala Ser Ile Leu Asn Leu Cys Ile Ile Ser Val Asp Arg Tyr Trp Ala
 130 135 140

 Ile Ser Ser Pro Phe Arg Tyr Glu Arg Lys Met Thr Gln Arg Val Ala
 145 150 155 160

 Phe Val Met Ile Ser Val Thr Trp Thr Leu Ser Val Leu Ile Ser Phe
 165 170 175

 Ile Pro Val Gln Leu Asn Trp His Lys Ala Ser Asp Glu Glu Val Trp
 180 185 190

 Ile Asn Gly Thr Ser Phe Gly Glu Lys Ser Glu Asn Cys Asp Ser Ser
 195 200 205

 Leu Asn Arg Glu Tyr Ala Ile Ser Ser Ser Leu Ile Ser Phe Tyr Ile
 210 215 220

 Pro Val Ala Ile Met Ile Val Thr Tyr Thr Arg Ile Tyr Arg Ile Ala
 225 230 235 240

 Gln Ile Gln Ile Arg Arg Ile Ser Ser Leu Glu Arg Ala Ala Glu His
 245 250 255

 Ala Gln Ser Cys Arg Thr Asn Arg Leu Glu Cys Gln His His Asn Thr
 260 265 270

 Leu Lys Thr Ser Ile Lys Arg Glu Thr Lys Val Phe Lys Thr Leu Ser
 275 280 285

 Val Ile Met Gly Val Phe Val Cys Cys Trp Leu Pro Phe Phe Ile Leu
 290 295 300

 Asn Cys Ile Val Pro Phe Cys Asp Arg Pro Pro Thr Asp His Thr Ala
 305 310 315 320

 Gly Leu Pro Cys Val Ser Asp Thr Thr Phe Asp Val Phe Val Trp Phe
 325 330 335

Gly Trp Thr Asn Ser Ser Leu Asn Pro Ile Ile Tyr Ala Phe Asn Ala
 340 345 350

 Asp Phe Arg Lys Ala Phe Ala Ser Leu Leu Gly Cys Arg Asn Phe Cys
 355 360 365

 Ser Arg Thr Pro Val Glu Thr Val Asn Ile Ser Asn Glu Leu Val Ser
 370 375 380

 Tyr Asn Gln Asp Thr Leu Phe His Lys Glu Ile Val Thr Ala Tyr Val
 385 390 395 400

 Asn Met Ile Pro Asn Val Val Asp Cys Ile Asp Asp Asn Glu Asp Thr
 405 410 415

 Phe Asp Arg Ile Ser Gln Phe Ser His Asn Asn Glu Ile Ala Thr Asp
 420 425 430

 Ser Val Cys Asp Leu Asp Asp Cys Glu Ala Asp Ile Cys Leu Asp Arg
 435 440 445

 Leu Ala Pro Phe Thr Pro Asn Gly Leu His
 450 455

<210> 52
 <211> 879
 <212> PRT
 <213> Rattus norvegicus

<400> 52
 Met Lys Met Leu Thr Arg Leu Gln Ile Leu Met Leu Ala Leu Phe Ser
 1 5 10 15

 Lys Gly Phe Leu Leu Ser Leu Gly Asp His Asn Phe Met Arg Arg Glu
 20 25 30

 Ile Lys Ile Glu Gly Asp Leu Val Leu Gly Gly Leu Phe Pro Ile Asn
 35 40 45

 Glu Lys Gly Thr Gly Thr Glu Glu Cys Gly Arg Ile Asn Glu Asp Arg
 50 55 60

 Gly Ile Gln Arg Leu Glu Ala Met Leu Phe Ala Ile Asp Glu Ile Asn
 65 70 75 80

 Lys Asp Asn Tyr Leu Leu Pro Gly Val Lys Leu Gly Val His Ile Leu
 85 90 95

 Asp Thr Cys Ser Arg Asp Thr Tyr Ala Leu Glu Gln Ser Leu Glu Phe
 100 105 110

 Val Arg Ala Ser Leu Thr Lys Val Asp Glu Ala Glu Tyr Met Cys Pro
 115 120 125

 Asp Gly Ser Tyr Ala Ile Gln Glu Asn Ile Pro Leu Leu Ile Ala Gly
 130 135 140

Val Ile Gly Gly Ser Tyr Ser Ser Val Ser Ile Gln Val Ala Asn Leu
145 150 155 160

Leu Arg Leu Phe Gln Ile Pro Gln Ile Ser Tyr Ala Ser Thr Ser Ala
165 170 175

Lys Leu Ser Asp Lys Ser Arg Tyr Asp Tyr Phe Ala Arg Thr Val Pro
180 185 190

Pro Asp Phe Tyr Gln Ala Lys Ala Met Ala Glu Ile Leu Arg Phe Phe
195 200 205

Asn Trp Thr Tyr Val Ser Thr Val Ala Ser Glu Gly Asp Tyr Gly Glu
210 215 220

Thr Gly Ile Glu Ala Phe Glu Gln Glu Ala Arg Leu Arg Asn Ile Cys
225 230 235 240

Ile Ala Thr Ala Glu Lys Val Gly Arg Ser Asn Ile Arg Lys Ser Tyr
245 250 255

Asp Ser Val Ile Arg Glu Leu Leu Gln Lys Pro Asn Ala Arg Val Val
260 265 270

Val Leu Phe Met Arg Ser Asp Asp Ser Arg Glu Leu Ile Ala Ala Ala
275 280 285

Asn Arg Val Asn Ala Ser Phe Thr Trp Val Ala Ser Asp Gly Trp Gly
290 295 300

Ala Gln Glu Ser Ile Val Lys Gly Ser Glu His Val Ala Tyr Gly Ala
305 310 315 320

Ile Thr Leu Glu Leu Ala Ser His Pro Val Arg Gln Phe Asp Arg Tyr
325 330 335

Phe Gln Ser Leu Asn Pro Tyr Asn Asn His Arg Asn Pro Trp Phe Arg
340 345 350

Asp Phe Trp Glu Gln Lys Phe Gln Cys Ser Leu Gln Asn Lys Arg Asn
355 360 365

His Arg Gln Val Cys Asp Lys His Leu Ala Ile Asp Ser Ser Asn Tyr
370 375 380

Glu Gln Glu Ser Lys Ile Met Phe Val Val Asn Ala Val Tyr Ala Met
385 390 395 400

Ala His Ala Leu His Lys Met Gln Arg Thr Leu Cys Pro Asn Thr Thr
405 410 415

Lys Leu Cys Asp Ala Met Lys Ile Leu Asp Gly Lys Lys Leu Tyr Lys
420 425 430

Glu Tyr Leu Leu Lys Ile Asn Phe Thr Ala Pro Phe Asn Pro Asn Lys
435 440 445

Gly Ala Asp Ser Ile Val Lys Phe Asp Thr Phe Gly Asp Gly Met Gly
450 455 460

Arg Tyr Asn Val Phe Asn Leu Gln Gln Thr Gly Gly Lys Tyr Ser Tyr
465 470 475 480

Leu Lys Val Gly His Trp Ala Glu Thr Leu Ser Leu Asp Val Asp Ser
485 490 495

Ile His Trp Ser Arg Asn Ser Val Pro Thr Ser Gln Cys Ser Asp Pro
500 505 510

Cys Ala Pro Asn Glu Met Lys Asn Met Gln Pro Gly Asp Val Cys Cys
515 520 525

Trp Ile Cys Ile Pro Cys Glu Pro Tyr Glu Tyr Leu Val Asp Glu Phe
530 535 540

Thr Cys Met Asp Cys Gly Pro Gly Gln Trp Pro Thr Ala Asp Leu Ser
545 550 555 560

Gly Cys Tyr Asn Leu Pro Glu Asp Tyr Ile Lys Trp Glu Asp Ala Trp
565 570 575

Ala Ile Gly Pro Val Thr Ile Ala Cys Leu Gly Phe Leu Cys Thr Cys
580 585 590

Ile Val Ile Thr Val Phe Ile Lys His Asn Asn Thr Pro Leu Val Lys
595 600 605

Ala Ser Gly Arg Glu Leu Cys Tyr Ile Leu Leu Phe Gly Val Ser Leu
610 615 620

Ser Tyr Cys Met Thr Phe Phe Ile Ala Lys Pro Ser Pro Val Ile
625 630 635 640

Cys Ala Leu Arg Arg Leu Gly Leu Gly Thr Ser Phe Ala Ile Cys Tyr
645 650 655

Ser Ala Leu Leu Thr Lys Thr Asn Cys Ile Ala Arg Ile Phe Asp Gly
660 665 670

Val Lys Asn Gly Ala Gln Arg Pro Lys Phe Ile Ser Pro Ser Ser Gln
675 680 685

Val Phe Ile Cys Leu Gly Leu Ile Leu Val Gln Ile Val Met Val Ser
690 695 700

Val Trp Leu Ile Leu Glu Thr Pro Gly Thr Arg Arg Tyr Thr Leu Pro
705 710 715 720

Glu Lys Arg Glu Thr Val Ile Leu Lys Cys Asn Val Lys Asp Ser Ser
725 730 735

Met Leu Ile Ser Leu Thr Tyr Asp Val Val Leu Val Ile Leu Cys Thr
740 745 750

Val Tyr Ala Phe Lys Thr Arg Lys Cys Pro Glu Asn Phe Asn Glu Ala
 755 760 765
 Lys Phe Ile Gly Phe Thr Met Tyr Thr Thr Cys Ile Ile Trp Leu Ala
 770 775 780
 Phe Leu Pro Ile Phe Tyr Val Thr Ser Ser Asp Tyr Arg Val Gln Thr
 785 790 795 800
 Thr Thr Met Cys Ile Ser Val Ser Leu Ser Gly Phe Val Val Leu Gly
 805 810 815
 Cys Leu Phe Ala Pro Lys Val His Ile Val Leu Phe Gln Pro Gln Lys
 820 825 830
 Asn Val Val Thr His Arg Leu His Leu Asn Arg Phe Ser Val Ser Gly
 835 840 845
 Thr Ala Thr Thr Tyr Ser Gln Ser Ser Ala Ser Thr Tyr Val Pro Thr
 850 855 860
 Val Cys Asn Gly Arg Glu Val Leu Asp Ser Thr Thr Ser Ser Leu
 865 870 875

<210> 53
 <211> 879
 <212> PRT
 <213> Mus musculus

<400> 53
 Met Lys Met Leu Thr Arg Leu Gln Val Leu Met Leu Ala Leu Phe Ser
 1 5 10 15
 Lys Gly Phe Leu Val Ser Leu Gly Asp His Asn Phe Met Arg Arg Glu
 20 25 30
 Ile Lys Ile Glu Gly Asp Leu Val Leu Gly Gly Leu Phe Pro Ile Asn
 35 40 45
 Glu Lys Gly Thr Gly Thr Glu Glu Cys Gly Arg Ile Asn Glu Asp Arg
 50 55 60
 Gly Ile Gln Arg Leu Glu Ala Met Leu Phe Ala Ile Asp Glu Ile Asn
 65 70 75 80
 Lys Asp Asn Tyr Leu Leu Pro Gly Val Lys Leu Gly Val His Ile Leu
 85 90 95
 Asp Thr Cys Ser Arg Asp Thr Tyr Ala Leu Glu Gln Ser Leu Glu Phe
 100 105 110
 Val Arg Ala Ser Leu Thr Lys Val Asp Glu Ala Glu Tyr Met Cys Pro
 115 120 125
 Asp Gly Ser Tyr Ala Ile Gln Glu Asn Ile Pro Leu Leu Ile Ala Gly

130	135	140
Val Ile Gly Gly Ser Tyr Ser Ser Val Ser Ile Gln Val Ala Asn Leu		
145	150	155
Leu Arg Leu Phe Gln Ile Pro Gln Ile Ser Tyr Ala Ser Thr Ser Ala		
165	170	175
Lys Leu Ser Asp Lys Ser Arg Tyr Asp Tyr Phe Ala Arg Thr Val Pro		
180	185	190
Pro Asp Phe Tyr Gln Ala Lys Ala Met Ala Glu Ile Leu Arg Tyr Phe		
195	200	205
Asn Trp Thr Tyr Val Ser Thr Val Ala Ser Glu Gly Asp Tyr Gly Glu		
210	215	220
Thr Gly Ile Glu Ala Phe Glu Gln Glu Ala Arg Leu Arg Asn Ile Cys		
225	230	235
Ile Ala Thr Ala Glu Lys Val Gly Arg Ser Asn Ile Arg Lys Ser Tyr		
245	250	255
Asp Ser Val Ile Arg Glu Leu Leu Gln Lys Pro Asn Ala Arg Val Val		
260	265	270
Val Leu Phe Met Arg Ser Asp Asp Ser Arg Glu Leu Ile Ala Ala Ala		
275	280	285
Ser Arg Val Asn Ala Ser Phe Thr Trp Val Ala Ser Asp Gly Trp Gly		
290	295	300
Ala Gln Glu Ser Ile Val Lys Gly Ser Glu His Val Ala Tyr Gly Ala		
305	310	315
Ile Thr Leu Glu Leu Ala Ser His Pro Val Arg Gln Phe Asp Arg Tyr		
325	330	335
Phe Gln Ser Leu Asn Pro Tyr Asn Asn His Arg Asn Pro Trp Phe Arg		
340	345	350
Asp Phe Trp Glu Gln Lys Phe Gln Cys Ser Leu Gln Asn Lys Arg Asn		
355	360	365
His Arg Gln Ile Cys Asp Lys His Leu Ala Ile Asp Ser Ser Asn Tyr		
370	375	380
Glu Gln Glu Ser Lys Ile Met Phe Val Val Asn Ala Val Tyr Ala Met		
385	390	395
Ala His Ala Leu His Lys Met Gln Arg Thr Leu Cys Pro Asn Thr Thr		
405	410	415
Lys Leu Cys Asp Ala Met Lys Ile Leu Asp Gly Lys Lys Leu Tyr Lys		
420	425	430
Asp Tyr Leu Leu Lys Ile Asn Phe Thr Ala Pro Phe Asn Pro Asn Lys		

435

440

445

Gly Ala Asp Ser Ile Val Lys Phe Asp Thr Tyr Gly Asp Gly Met Gly
 450 455 460

Arg Tyr Asn Val Phe Asn Phe Gln His Ile Gly Gly Lys Tyr Ser Tyr
 465 470 475 480

Leu Lys Val Gly His Trp Ala Glu Thr Leu Tyr Leu Asp Val Asp Ser
 485 490 495

Ile His Trp Ser Arg Asn Ser Val Pro Thr Ser Gln Cys Ser Asp Pro
 500 505 510

Cys Ala Pro Asn Glu Met Lys Asn Met Gln Pro Gly Asp Val Cys Cys
 515 520 525

Trp Ile Cys Ile Pro Cys Glu Pro Tyr Glu Tyr Leu Val Asp Glu Phe
 530 535 540

Thr Cys Met Asp Cys Gly Pro Gly Gln Trp Pro Thr Ala Asp Leu Ser
 545 550 555 560

Gly Cys Tyr Asn Leu Pro Glu Asp Tyr Ile Arg Trp Glu Asp Ala Trp
 565 570 575

Ala Ile Gly Pro Val Thr Ile Ala Cys Leu Gly Phe Met Cys Thr Cys
 580 585 590

Ile Val Ile Thr Val Phe Ile Lys His Asn Asn Thr Pro Leu Val Lys
 595 600 605

Ala Ser Gly Arg Glu Leu Cys Tyr Ile Leu Leu Phe Gly Val Ser Leu
 610 615 620

Ser Tyr Cys Met Thr Phe Phe Ile Ala Lys Pro Ser Pro Val Ile
 625 630 635 640

Cys Ala Leu Arg Arg Leu Gly Leu Gly Thr Ser Phe Ala Ile Cys Tyr
 645 650 655

Ser Ala Leu Leu Thr Lys Thr Asn Cys Ile Ala Arg Ile Phe Asp Gly
 660 665 670

Val Lys Asn Gly Ala Gln Arg Pro Lys Phe Ile Ser Pro Ser Ser Gln
 675 680 685

Val Phe Ile Cys Leu Gly Leu Ile Leu Val Gln Ile Val Met Val Ser
 690 695 700

Val Trp Leu Ile Leu Glu Thr Pro Gly Thr Arg Arg Tyr Thr Leu Pro
 705 710 715 720

Glu Lys Arg Glu Thr Val Ile Leu Lys Cys Asn Val Lys Asp Ser Ser
 725 730 735

Met Leu Ile Ser Leu Thr Tyr Asp Val Val Leu Val Ile Leu Cys Thr

740

745

750

Val Tyr Ala Phe Lys Thr Arg Lys Cys Pro Glu Asn Phe Asn Glu Ala
755 760 765

Lys Phe Ile Gly Phe Thr Met Tyr Thr Thr Cys Ile Ile Trp Leu Ala
770 775 780

Phe Leu Pro Ile Phe Tyr Val Thr Ser Ser Asp Tyr Arg Val Gln Thr
785 790 795 800

Thr Thr Met Cys Ile Ser Val Ser Leu Ser Gly Phe Val Val Leu Gly
805 810 815

Cys Leu Phe Ala Pro Lys Val His Ile Val Leu Phe Gln Pro Gln Lys
820 825 830

Asn Val Val Thr His Arg Leu His Leu Asn Arg Phe Ser Val Ser Gly
835 840 845

Thr Ala Thr Thr Tyr Ser Gln Ser Ser Ala Ser Thr Tyr Val Pro Thr
850 855 860

Val Cys Asn Gly Arg Glu Val Leu Asp Ser Thr Thr Ser Ser Leu
865 870 875

<210> 54

<211> 879

<212> PRT

<213> Mus musculus

<400> 54

Met Lys Met Leu Thr Arg Leu Gln Val Leu Met Leu Ala Leu Phe Ser
1 5 10 15

Lys Gly Phe Leu Val Ser Leu Gly Asp His Asn Phe Met Arg Arg Glu
20 25 30

Ile Lys Ile Glu Gly Asp Leu Val Leu Gly Gly Leu Phe Pro Ile Asn
35 40 45

Glu Lys Gly Thr Gly Thr Glu Glu Cys Arg Gly Ile Asn Glu Asp Arg
50 55 60

Gly Ile Gln Arg Leu Glu Ala Met Leu Phe Ala Ile Asp Glu Ile Asn
65 70 75 80

Lys Asp Asn Tyr Leu Leu Pro Gly Val Lys Leu Gly Val His Ile Leu
85 90 95

Asp Thr Cys Ser Arg Asp Thr Tyr Ala Leu Glu Gln Ser Leu Glu Phe
100 105 110

Val Arg Ala Ser Leu Thr Lys Val Asp Glu Ala Glu Tyr Met Cys Pro
115 120 125

Asp Gly Ser Tyr Ala Ile Gln Glu Asn Ile Pro Leu Leu Ile Ala Gly
130 135 140

Val Ile Gly Gly Ser Tyr Ser Ser Val Ser Ile Gln Val Ala Asn Leu
145 150 155 160

Leu Arg Leu Phe Gln Ile Pro Gln Ile Ser Tyr Ala Ser Thr Ser Ala
165 170 175

Lys Leu Ser Asp Lys Ser Arg Tyr Asp Tyr Phe Ala Arg Thr Val Pro
180 185 190

Pro Asp Phe Tyr Gln Ala Lys Ala Met Ala Glu Ile Leu Arg Tyr Phe
195 200 205

Asn Trp Thr Tyr Val Ser Thr Val Ala Ser Glu Gly Asp Tyr Gly Glu
210 215 220

Thr Gly Ile Glu Ala Phe Glu Gln Glu Ala Arg Leu Arg Asn Ile Cys
225 230 235 240

Ile Ala Thr Ala Glu Lys Val Gly Arg Ser Asn Ile Arg Lys Ser Tyr
245 250 255

Asp Ser Val Ile Arg Glu Leu Leu Gln Lys Pro Asn Ala Arg Val Val
260 265 270

Val Leu Phe Met Arg Ser Asp Asp Ser Arg Glu Leu Ile Ala Ala Ala
275 280 285

Ser Arg Val Asn Ala Ser Phe Thr Trp Val Ala Ser Asp Gly Trp Gly
290 295 300

Ala Gln Glu Ser Ile Val Lys Gly Ser Glu His Val Ala Tyr Gly Ala
305 310 315 320

Ile Thr Leu Glu Leu Ala Ser His Pro Val Arg Gln Phe Asp Arg Tyr
325 330 335

Phe Gln Ser Leu Asn Pro Tyr Asn Asn His Arg Asn Pro Trp Phe Arg
340 345 350

Asp Phe Trp Glu Gln Lys Phe Gln Cys Ser Leu Gln Asn Lys Arg Asn
355 360 365

His Arg Gln Ile Cys Asp Lys His Leu Ala Ile Asp Ser Ser Asn Tyr
370 375 380

Glu Gln Glu Ser Lys Ile Met Phe Val Val Asn Ala Val Tyr Ala Met
385 390 395 400

Ala His Ala Leu His Lys Met Gln Arg Thr Leu Cys Pro Asn Thr Thr
405 410 415

Lys Leu Cys Asp Ala Met Lys Ile Leu Asp Gly Lys Lys Leu Tyr Lys
420 425 430

Asp	Tyr	Leu	Leu	Lys	Ile	Asn	Phe	Thr	Ala	Pro	Phe	Asn	Pro	Asn	Lys
					435		440					445			
Gly	Ala	Asp	Ser	Ile	Val	Lys	Phe	Asp	Thr	Tyr	Gly	Asp	Gly	Met	Gly
					450		455				460				
Arg	Tyr	Asn	Val	Phe	Asn	Phe	Gln	His	Ile	Gly	Gly	Lys	Tyr	Ser	Tyr
					465		470			475			480		
Leu	Lys	Val	Gly	His	Trp	Ala	Glu	Thr	Leu	Tyr	Leu	Asp	Val	Asp	Ser
					485		490			495					
Ile	His	Trp	Ser	Arg	Asn	Ser	Val	Pro	Thr	Ser	Gln	Cys	Ser	Asp	Pro
					500		505				510				
Cys	Ala	Pro	Asn	Glu	Met	Lys	Asn	Met	Gln	Pro	Gly	Asp	Val	Cys	Cys
					515		520			525					
Trp	Ile	Cys	Ile	Pro	Cys	Glu	Pro	Tyr	Glu	Tyr	Leu	Val	Asp	Glu	Phe
					530		535			540					
Thr	Cys	Met	Asp	Cys	Gly	Pro	Gly	Gln	Trp	Pro	Thr	Ala	Asp	Leu	Ser
					545		550			555			560		
Gly	Cys	Tyr	Asn	Leu	Pro	Glu	Asp	Tyr	Ile	Arg	Trp	Glu	Asp	Ala	Trp
					565		570			575					
Ala	Ile	Gly	Pro	Val	Thr	Ile	Ala	Cys	Leu	Gly	Phe	Met	Cys	Thr	Cys
					580		585			590					
Ile	Val	Ile	Thr	Val	Phe	Ile	Lys	His	Asn	Asn	Thr	Pro	Leu	Val	Lys
					595		600			605					
Ala	Ser	Gly	Arg	Glu	Leu	Cys	Tyr	Ile	Leu	Leu	Phe	Gly	Val	Ser	Leu
					610		615			620					
Ser	Tyr	Cys	Met	Thr	Phe	Phe	Phe	Ile	Ala	Lys	Pro	Ser	Pro	Val	Ile
					625		630			635			640		
Cys	Ala	Leu	Arg	Arg	Leu	Gly	Leu	Gly	Thr	Ser	Phe	Ala	Ile	Cys	Tyr
					645		650			655					
Ser	Ala	Leu	Leu	Thr	Lys	Thr	Asn	Cys	Ile	Ala	Arg	Ile	Phe	Asp	Gly
					660		665			670					
Val	Lys	Asn	Gly	Ala	Gln	Arg	Pro	Lys	Phe	Ile	Ser	Pro	Ser	Ser	Gln
					675		680			685					
Val	Phe	Ile	Cys	Leu	Gly	Leu	Ile	Leu	Val	Gln	Ile	Val	Met	Val	Ser
					690		695			700					
Val	Trp	Leu	Ile	Leu	Glu	Thr	Pro	Gly	Thr	Arg	Arg	Tyr	Thr	Leu	Pro
					705		710			715			720		
Glu	Lys	Arg	Glu	Thr	Val	Ile	Leu	Lys	Cys	Asn	Val	Lys	Asp	Ser	Ser
					725		730			735					

Met Leu Ile Ser Leu Thr Tyr Asp Val Val Leu Val Ile Leu Cys Thr
740 745 750

Val Tyr Ala Phe Lys Thr Arg Lys Cys Pro Glu Asn Phe Asn Glu Ala
755 760 765

Lys Phe Ile Gly Phe Thr Met Tyr Thr Thr Cys Ile Ile Trp Leu Ala
770 775 780

Phe Leu Pro Ile Phe Tyr Val Thr Ser Ser Asp Tyr Arg Val Gln Thr
785 790 795 800

Thr Thr Met Cys Ile Ser Val Ser Leu Ser Gly Phe Val Val Leu Gly
805 810 815

Cys Leu Phe Ala Pro Lys Val His Ile Val Leu Phe Gln Pro Gln Lys
820 825 830

Asn Val Val Thr His Arg Leu His Leu Asn Arg Phe Ser Val Ser Gly
835 840 845

Thr Ala Thr Thr Tyr Ser Gln Ser Ser Ala Ser Thr Tyr Val Pro Thr
850 855 860

Val Cys Asn Gly Arg Glu Val Leu Asp Ser Thr Thr Ser Ser Leu
865 870 875

<210> 55

<211> 442

<212> PRT

<213> Homo sapiens

<400> 55

Met Gly Leu Ala Met Glu His Gly Gly Ser Tyr Ala Arg Ala Gly Gly
1 5 10 15

Ser Ser Arg Gly Cys Trp Tyr Tyr Leu Arg Tyr Phe Phe Leu Phe Val
20 25 30

Ser Leu Ile Gln Phe Leu Ile Ile Leu Gly Leu Val Leu Phe Met Val
35 40 45

Tyr Gly Asn Val His Val Ser Thr Glu Ser Asn Leu Gln Ala Thr Glu
50 55 60

Arg Arg Ala Glu Gly Leu Tyr Ser Gln Leu Leu Gly Leu Thr Ala Ser
65 70 75 80

Gln Ser Asn Leu Thr Lys Glu Leu Asn Phe Thr Thr Arg Ala Lys Asp
85 90 95

Ala Ile Met Gln Met Trp Leu Asn Ala Arg Arg Asp Leu Asp Arg Ile
100 105 110

Asn Ala Ser Phe Arg Gln Cys Gln Gly Asp Arg Val Ile Tyr Thr Asn
115 120 125

Asn His Arg Tyr Met Ala Ala Ile Ile Leu Ser Glu Lys Gln Cys Arg
 130 135 140

 Asp Gln Phe Lys Asp Met Asn Lys Ser Cys Asp Ala Leu Leu Phe Met
 145 150 155 160

 Leu Asn Gln Lys Val Lys Thr Leu Glu Val Glu Ile Ala Lys Glu Lys
 165 170 175

 Thr Ile Cys Thr Lys Asp Lys Glu Ser Val Leu Leu Asn Lys Arg Val
 180 185 190

 Ala Glu Glu Gln Leu Val Glu Cys Val Lys Thr Arg Glu Leu Gln His
 195 200 205

 Gln Glu Arg Gln Leu Ala Lys Glu Gln Leu Gln Lys Val Gln Ala Leu
 210 215 220

 Cys Leu Pro Leu Asp Lys Asp Lys Phe Glu Met Asp Leu Arg Asn Leu
 225 230 235 240

 Trp Arg Asp Ser Ile Ile Pro Arg Ser Leu Asp Asn Leu Gly Tyr Asn
 245 250 255

 Leu Tyr His Pro Leu Gly Ser Glu Leu Ala Ser Ile Arg Arg Ala Cys
 260 265 270

 Asp His Met Pro Ser Leu Met Ser Ser Lys Val Glu Glu Leu Ala Arg
 275 280 285

 Ser Leu Arg Ala Asp Ile Glu Arg Val Ala Arg Glu Asn Ser Asp Leu
 290 295 300

 Gln Arg Gln Lys Leu Glu Ala Gln Gln Gly Leu Arg Ala Ser Gln Glu
 305 310 315 320

 Ala Lys Gln Lys Val Glu Lys Glu Ala Gln Ala Arg Glu Ala Lys Leu
 325 330 335

 Gln Ala Glu Cys Ser Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala
 340 345 350

 Val Leu Arg Lys Glu Arg Asp Asn Leu Ala Lys Glu Leu Glu Glu Lys
 355 360 365

 Lys Arg Glu Ala Glu Gln Leu Arg Met Glu Leu Ala Ile Arg Asn Ser
 370 375 380

 Ala Leu Asp Thr Cys Ile Lys Thr Lys Ser Gln Pro Met Met Pro Val
 385 390 395 400

 Ser Arg Pro Met Gly Pro Val Pro Asn Pro Gln Pro Ile Asp Pro Ala
 405 410 415

 Ser Leu Glu Glu Phe Lys Arg Lys Ile Leu Glu Ser Gln Arg Pro Pro
 420 425 430

Ala Gly Ile Pro Val Ala Pro Ser Ser Gly
435 440

<210> 56
<211> 442
<212> PRT
<213> Homo sapiens

<400> 56
Met Gly Leu Ala Met Glu His Gly Gly Ser Tyr Ala Arg Ala Gly Gly
1 5 10 15

Ser Ser Arg Gly Cys Trp Tyr Tyr Leu Arg Tyr Phe Phe Leu Phe Val
20 25 30

Ser Leu Ile Gln Phe Leu Ile Ile Leu Gly Leu Val Leu Phe Met Val
35 40 45

Tyr Gly Asn Val His Val Ser Thr Glu Ser Asn Leu Gln Ala Thr Glu
50 55 60

Arg Arg Ala Glu Gly Leu Tyr Ser Gln Leu Leu Gly Leu Thr Ala Ser
65 70 75 80

Gln Ser Asn Leu Thr Lys Glu Leu Asn Phe Thr Thr Arg Ala Lys Asp
85 90 95

Ala Ile Met Gln Met Trp Leu Asn Ala Arg Arg Asp Leu Asp Arg Ile
100 105 110

Asn Ala Ser Phe Arg Gln Cys Gln Gly Asp Arg Val Ile Tyr Thr Asn
115 120 125

Asn Gln Arg Tyr Met Ala Ala Ile Ile Leu Ser Glu Lys Gln Cys Arg
130 135 140

Asp Gln Phe Lys Asp Met Asn Lys Ser Cys Asp Ala Leu Leu Phe Met
145 150 155 160

Leu Asn Gln Lys Val Lys Thr Leu Glu Val Glu Ile Ala Lys Glu Lys
165 170 175

Thr Ile Cys Thr Lys Asp Lys Glu Ser Val Leu Leu Asn Lys Arg Val
180 185 190

Ala Glu Glu Gln Leu Val Glu Cys Val Lys Thr Arg Glu Leu Gln His
195 200 205

Gln Glu Arg Gln Leu Ala Lys Glu Gln Leu Gln Lys Val Gln Ala Leu
210 215 220

Cys Leu Pro Leu Asp Lys Asp Lys Phe Glu Met Asp Leu Arg Asn Leu
225 230 235 240

Trp Arg Asp Ser Ile Ile Pro Arg Ser Leu Asp Asn Leu Gly Tyr Asn

245

250

255

Leu Tyr His Pro Leu Gly Ser Glu Leu Ala Ser Ile Arg Arg Ala Cys
 260 265 270

Asp His Met Pro Ser Leu Met Ser Ser Lys Val Glu Glu Leu Ala Arg
 275 280 285

Ser Leu Arg Ala Asp Ile Glu Arg Val Ala Arg Glu Asn Ser Asp Leu
 290 295 300

Gln Arg Gln Lys Leu Glu Ala Gln Gln Gly Leu Arg Ala Ser Gln Glu
 305 310 315 320

Ala Lys Gln Lys Val Glu Lys Glu Ala Gln Ala Arg Glu Ala Lys Leu
 325 330 335

Gln Ala Glu Cys Ser Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala
 340 345 350

Val Leu Arg Lys Glu Arg Asp Asn Leu Ala Lys Glu Leu Glu Glu Lys
 355 360 365

Lys Arg Glu Ala Glu Gln Leu Arg Met Glu Leu Ala Ile Arg Asn Ser
 370 375 380

Ala Leu Asp Thr Cys Ile Lys Thr Lys Ser Gln Pro Met Met Pro Val
 385 390 395 400

Ser Arg Pro Met Gly Pro Val Pro Asn Pro Gln Pro Ile Asp Pro Ala
 405 410 415

Ser Leu Glu Glu Phe Lys Arg Lys Ile Leu Glu Ser Gln Arg Pro Pro
 420 425 430

Ala Gly Ile Pro Val Ala Pro Ser Ser Gly
 435 440

<210> 57

<211> 438

<212> PRT

<213> Rattus norvegicus

<400> 57

Met Gly Leu Ser Met Asp Arg Ser Pro Tyr Ser Arg Thr Gly Asp Arg
 1 5 10 15

Asp Arg Gly Cys Trp Tyr Tyr Leu Arg Tyr Phe Phe Leu Phe Val Ser
 20 25 30

Leu Ile Gln Phe Leu Ile Ile Leu Gly Leu Val Leu Phe Met Ile Tyr
 35 40 45

Gly Asn Val His Ala Thr Thr Glu Ser Ser Leu Arg Ala Thr Glu Ile
 50 55 60

Arg	Ala	Asp	Asn	Leu	Tyr	Ser	Gln	Val	Val	Gly	Leu	Ser	Ala	Ala	Gln
65				70				75						80	
Ala	Asn	Leu	Ser	Lys	Gln	Leu	Asn	Ile	Ser	Thr	Leu	Val	Lys	Asp	Thr
				85				90					95		
Val	Met	Gln	Gln	Leu	Leu	Thr	Thr	Arg	Arg	Glu	Val	Glu	Arg	Ile	Asn
				100				105					110		
Ala	Ser	Phe	Arg	Gln	Cys	Gln	Gly	Asp	Leu	Ile	Thr	Tyr	Ile	Asn	Tyr
				115				120				125			
Asn	Arg	Phe	Ile	Ala	Ala	Ile	Ile	Leu	Ser	Glu	Lys	Gln	Cys	Gln	Glu
				130				135			140				
Gln	Leu	Lys	Glu	Gly	Asn	Lys	Thr	Cys	Glu	Ala	Leu	Leu	Phe	Lys	Leu
				145				150			155		160		
Gly	Glu	Lys	Val	Lys	Thr	Leu	Glu	Met	Glu	Val	Val	Lys	Glu	Lys	Ala
				165				170				175			
Val	Cys	Ser	Lys	Asp	Lys	Asp	Ser	Leu	Leu	Ala	Gly	Lys	Arg	Gln	Ala
				180				185				190			
Glu	Met	Gln	Gln	Glu	Ala	Cys	Gly	Lys	Ala	Arg	Glu	Gln	Gln	Lys	Gln
				195				200			205				
Asp	Gln	Gln	Val	Thr	Glu	Glu	Gln	Leu	Arg	Lys	Val	Gln	Ser	Leu	Cys
				210				215			220				
Leu	Pro	Leu	Asp	Gln	Glu	Lys	Phe	Gln	Ala	Asp	Val	Leu	Asn	Val	Trp
				225				230			235		240		
Arg	Asp	Ser	Leu	Val	Tyr	Arg	Ser	Leu	Asp	Asn	Ile	Gly	Tyr	His	Tyr
				245				250			255				
Ser	Leu	Met	Pro	Glu	Phe	Ser	Ser	Leu	Arg	Arg	Thr	Cys	Glu	Ser	Leu
				260				265			270				
Pro	Gly	Ile	Met	Thr	Thr	Lys	Val	Glu	Glu	Leu	Ala	Arg	Gly	Leu	Arg
				275				280			285				
Ala	Gly	Ile	Glu	Arg	Val	Thr	Arg	Glu	Asn	Gly	Glu	Leu	Arg	Arg	Gln
				290				295			300				
Lys	Leu	Glu	Leu	Glu	Arg	Ala	Ile	Gln	Gly	Glu	Arg	Glu	Ala	Arg	Thr
				305				310			315		320		
Arg	Ala	Gly	Thr	Glu	Ala	Gln	Ala	Arg	Glu	Thr	Gln	Leu	Arg	Thr	Glu
				325				330			335				
Cys	Ala	Arg	Gln	Thr	Gln	Leu	Ala	Leu	Glu	Glu	Lys	Ala	Ala	Leu	Arg
				340				345			350				
Thr	Gln	Arg	Asp	Asp	Leu	Glu	Arg	Gln	Leu	Glu	Ala	Arg	Lys	Arg	Glu
				355				360			365				

Leu Glu Gln Leu Arg Thr Glu Val Asp Val Arg Ile Ser Ala Leu Asp
370 375 380

Thr Cys Val Lys Ala Lys Ser Leu Pro Ala Ile Gln Pro Arg Leu Pro
385 390 395 400

Gly Pro Pro Pro Asn Pro Pro Pro Ile Asp Pro Ala Ser Leu Glu Glu
405 410 415

Phe Lys Lys Arg Ile Leu Glu Ser Gln Arg Pro Pro Leu Val Asn Pro
420 425 430

Ala Val Pro Pro Ser Gly
435

<210> 58

<211> 438

<212> PRT

<213> Mus musculus

<400> 58

Met Gly Leu Ser Met Asp Arg Ser Pro Tyr Ala Arg Thr Gly Asp Gln
1 5 10 15

Gln Arg Gly Cys Trp Tyr Tyr Leu Arg Tyr Phe Phe Leu Phe Val Ser
20 25 30

Leu Ile Gln Phe Leu Ile Ile Leu Gly Leu Val Leu Phe Met Ile Tyr
35 40 45

Gly Asn Val His Ala Thr Thr Glu Ser Ser Leu Arg Ala Thr Glu Ile
50 55 60

Arg Ala Asp Ser Leu Tyr Ser Gln Val Val Gly Leu Ser Ala Ser Gln
65 70 75 80

Ala Asn Leu Ser Lys Gln Leu Asn Ile Ser Leu Leu Val Lys Glu Thr
85 90 95

Val Met Gln Gln Leu Leu Thr Thr Arg Arg Glu Met Glu Arg Ile Asn
100 105 110

Ala Ser Phe Arg Gln Cys Gln Gly Asp Leu Ile Thr Tyr Ile Asn Tyr
115 120 125

Asn Arg Phe Ile Ala Ala Ile Ile Leu Ser Glu Lys Gln Cys Gln Glu
130 135 140

Gln Leu Lys Glu Val Asn Lys Thr Cys Glu Ala Leu Leu Phe Lys Leu
145 150 155 160

Gly Glu Lys Val Lys Thr Leu Glu Met Glu Val Ala Lys Glu Lys Ala
165 170 175

Val Cys Ser Lys Asp Lys Glu Ser Leu Leu Ala Gly Lys Arg Gln Thr
180 185 190

Glu Glu Gln Leu Glu Ala Cys Gly Lys Ala Arg Glu Arg Gln Gln Gln
 195 200 205

Glu Gln Gln Val Thr Glu Glu Asn Leu Arg Lys Val Gln Ser Leu Cys
 210 215 220

Ile Pro Leu Asp Gln Glu Lys Phe Gln Ala Asp Val Leu Ser Ala Trp
 225 230 235 240

Arg Asp Ser Leu Ile Tyr Arg Thr Leu Glu Thr Leu Pro Tyr His Tyr
 245 250 255

Gln Leu Met Pro Glu Tyr Ala Ser Leu Arg Arg Thr Cys Glu Ser Leu
 260 265 270

Pro Gly Ile Met Thr Thr Lys Ile Glu Glu Leu Ala Arg Gly Leu Arg
 275 280 285

Ala Gly Ile Glu Arg Val Thr Arg Glu Asn Ala Glu Leu Arg Arg Gln
 290 295 300

Lys Leu Glu Leu Glu Arg Ala Ala Gln Ala Ala Gln Glu Ala Arg Ala
 305 310 315 320

Arg Ala Gly Thr Glu Ala Gln Ala Arg Glu Thr Gln Leu Arg Ala Glu
 325 330 335

Cys Ala Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala Ala Leu Arg
 340 345 350

Ala Gln Arg Asp Asn Leu Glu Arg Glu Leu Glu Ala Arg Lys Arg Glu
 355 360 365

Leu Glu Gln Leu Arg Thr Glu Val Asp Val Arg Ile Ser Ala Leu Asp
 370 375 380

Thr Cys Val Lys Ala Lys Ser Leu Pro Ala Val Pro Pro Arg Val Ser
 385 390 395 400

Gly Pro Pro Pro Asn Pro Pro Pro Ile Asp Pro Ala Ser Leu Glu Glu
 405 410 415

Phe Lys Lys Arg Ile Leu Glu Ser Gln Arg Leu Pro Val Val Asn Pro
 420 425 430

Ala Ala Gln Pro Ser Gly
 435

<210> 59
 <211> 438
 <212> PRT
 <213> Mus musculus

<400> 59
 Met Gly Leu Ser Met Asp Arg Ser Pro Tyr Ala Arg Thr Gly Asp Gln

1	5	10	15												
Gln	Arg	Gly	Cys	Trp	Tyr	Tyr	Leu	Arg	Tyr	Phe	Phe	Leu	Phe	Val	Ser
				20			25							30	
Leu	Ile	Gln	Phe	Leu	Ile	Ile	Leu	Gly	Leu	Val	Leu	Phe	Met	Ile	Tyr
					35			40					45		
Gly	Asn	Val	His	Ala	Thr	Thr	Glu	Ser	Ser	Leu	Arg	Ala	Thr	Glu	Ile
					50		55				60				
Arg	Ala	Asp	Ser	Leu	Tyr	Ser	Gln	Val	Val	Gly	Leu	Ser	Ala	Ser	Gln
					65		70		75				80		
Ala	Asn	Leu	Ser	Lys	Gln	Leu	Asn	Ile	Ser	Leu	Leu	Val	Lys	Glu	Thr
					85			90					95		
Val	Met	Gln	Gln	Leu	Leu	Thr	Thr	Arg	Arg	Glu	Met	Glu	Arg	Ile	Asn
					100			105				110			
Ala	Ser	Phe	Arg	Gln	Cys	Gln	Gly	Asp	Leu	Ile	Thr	Tyr	Ile	Asn	Tyr
					115		120				125				
Asn	Arg	Phe	Ile	Ala	Ala	Ile	Ile	Leu	Ser	Glu	Lys	Gln	Cys	Gln	Glu
					130		135				140				
Gln	Leu	Lys	Glu	Val	Asn	Lys	Thr	Cys	Glu	Ala	Leu	Leu	Phe	Lys	Leu
					145		150				155		160		
Gly	Glu	Lys	Val	Lys	Thr	Leu	Glu	Met	Glu	Val	Ala	Lys	Glu	Lys	Ala
					165			170				175			
Val	Cys	Ser	Lys	Asp	Lys	Glu	Ser	Leu	Leu	Ala	Gly	Lys	Arg	Gln	Ala
					180			185				190			
Glu	Glu	Gln	Leu	Glu	Ala	Cys	Gly	Lys	Ala	Arg	Glu	Arg	Gln	Gln	
					195		200				205				
Glu	Gln	Gln	Val	Thr	Glu	Glu	Asn	Leu	Arg	Lys	Val	Gln	Ser	Leu	Cys
					210		215				220				
Ile	Pro	Leu	Asp	Gln	Glu	Lys	Phe	Gln	Ala	Asp	Val	Leu	Ser	Ala	Trp
					225		230				235		240		
Arg	Asp	Ser	Leu	Ile	Tyr	Arg	Thr	Leu	Glu	Thr	Leu	Pro	Tyr	His	Tyr
					245			250				255			
Gln	Leu	Met	Pro	Glu	Tyr	Ala	Ser	Leu	Arg	Arg	Thr	Cys	Glu	Ser	Leu
					260			265				270			
Pro	Gly	Ile	Met	Pro	Pro	Lys	Ile	Glu	Glu	Met	Ala	Arg	Gly	Val	Arg
					275		280				285				
Ala	Gly	Ile	Glu	Arg	Val	Thr	Arg	Glu	Asn	Ala	Glu	Leu	Arg	Arg	Gln
					290		295				300				
Lys	Leu	Glu	Leu	Glu	Arg	Ala	Ala	Gln	Arg	Ala	Gln	Glu	Ala	Arg	Ala

305	310	315	320
Arg Ala Gly Thr Glu Ala Gln Ala Arg Glu Thr Gln Leu Arg Ala Glu			
325	330	335	
Cys Ala Arg Gln Thr Gln Leu Ala Leu Glu Glu Lys Ala Ala Leu Arg			
340	345	350	
Ala Gln Arg Asp Asn Leu Glu Arg Glu Leu Glu Ala Arg Lys Arg Glu			
355	360	365	
Leu Glu Gln Leu Arg Thr Glu Val Asp Val Arg Ile Ser Ala Leu Asp			
370	375	380	
Thr Cys Val Lys Ala Lys Ser Leu Pro Ala Val Pro Pro Arg Val Ser			
385	390	395	400
Gly Pro Pro Pro Asn Pro Pro Pro Ile Asp Pro Ala Ser Leu Glu Glu			
405	410	415	
Phe Lys Lys Arg Ile Leu Glu Ser Gln Arg Leu Pro Val Val Asn Pro			
420	425	430	
Ala Ala Gln Pro Ser Gly			
435			

<210> 60
 <211> 1788
 <212> PRT
 <213> Rattus norvegicus

<400> 60			
Met Leu Arg Pro Met Gln Pro Thr Ser Arg Thr Leu Gln Arg Pro Pro			
1	5	10	15
Arg Gly Ala Leu Glu Ala Gly Gly Arg Arg Asn Cys Gln Asp Gln Val			
20	25	30	
Ala His Pro Asn Trp Asn Thr Gln Ser Val Gln Thr Pro Arg Val Arg			
35	40	45	
Arg Thr Leu Gly Ala Pro Val Pro Pro Ser Arg Lys Val Lys Ala Trp			
50	55	60	
Ala Pro Gly Thr Asp Gln Trp Pro Gly Val Ser Pro His Cys Lys Arg			
65	70	75	80
Ser Glu Ala Glu Ala Lys Pro Ser Gly Ser Gln Thr Val Asn Leu Thr			
85	90	95	
Gly Arg Ala Asn Asp Pro Cys Asp Leu Asp Ser Arg Val Gln Ala Thr			
100	105	110	
Ser Val Lys Val Thr Val Ala Gly Phe Gln Pro Gly Gly Ala Val Glu			
115	120	125	

Lys Leu Cys Gln Glu Ser Leu Gly Lys Leu Thr Thr Gly Asp Ala Cys
 130 135 140
 Val Ser Thr Ser Cys Glu Leu Ala Ser Ala Leu Ser His Leu Asp Ala
 145 150 155 160
 Ser His Leu Thr Glu Asn Leu Pro Lys Ala Ala Ser Glu Leu Gly Gln
 165 170 175
 Gln Pro Met Thr Ser Ser Asp Leu Ile Ser Ser Pro Gly Lys Lys Gly
 180 185 190
 Ala Ala His Pro Asp Pro Ser Lys Thr Ser Val Asp Thr Gly Gln Val
 195 200 205
 Ser Arg Pro Glu Asn Pro Ser Gln Pro Ala Ser Pro Arg Val Thr Lys
 210 215 220
 Cys Lys Ala Arg Ser Pro Val Arg Leu Pro His Glu Gly Ser Pro Ser
 225 230 235 240
 Pro Gly Glu Lys Ala Ala Ala Pro Pro Asp Tyr Ser Lys Thr Arg Ser
 245 250 255
 Ala Ser Glu Thr Ser Thr Pro His Asn Thr Arg Arg Val Ala Ala Leu
 260 265 270
 Arg Gly Ala Gly Pro Gly Ala Glu Gly Met Thr Pro Ala Gly Ala Val
 275 280 285
 Leu Pro Gly Asp Pro Leu Thr Ser Gln Glu Gln Arg Gln Gly Ala Pro
 290 295 300
 Gly Asn His Ser Lys Ala Leu Glu Met Thr Gly Ile His Ala Pro Glu
 305 310 315 320
 Ser Ser Gln Glu Pro Ser Leu Leu Glu Gly Ala Asp Ser Val Ser Ser
 325 330 335
 Arg Ala Pro Gln Ala Ser Leu Ser Met Leu Pro Ser Thr Asp Asn Thr
 340 345 350
 Lys Glu Ala Cys Gly His Val Ser Gly His Cys Cys Pro Gly Gly Ser
 355 360 365
 Arg Glu Ser Pro Val Thr Asp Ile Asp Ser Phe Ile Lys Glu Leu Asp
 370 375 380
 Ala Ser Ala Ala Arg Ser Pro Ser Ser Gln Thr Gly Asp Ser Gly Ser
 385 390 395 400
 Gln Glu Gly Ser Ala Gln Gly His Pro Pro Ala Gly Ala Gly Gly
 405 410 415
 Ser Ser Cys Arg Ala Glu Pro Val Pro Gly Gly Gln Thr Ser Ser Pro
 420 425 430

Arg Arg Ala Trp Ala Ala Gly Ala Pro Ala Tyr Pro Gln Trp Ala Ser
435 440 445

Gln Pro Ser Val Leu Asp Ser Ile Asn Pro Asp Lys His Phe Thr Val
450 455 460

Asn Lys Asn Phe Leu Ser Asn Tyr Ser Arg Asn Phe Ser Ser Phe His
465 470 475 480

Glu Asp Ser Thr Ser Leu Ser Gly Leu Gly Asp Ser Thr Glu Pro Ser
485 490 495

Leu Ser Ser Met Tyr Gly Asp Ala Glu Asp Ser Ser Asp Pro Glu
500 505 510

Ser Leu Thr Glu Ala Pro Arg Ala Ser Ala Arg Asp Gly Trp Ser Pro
515 520 525

Pro Arg Ser Arg Val Ser Leu His Lys Glu Asp Pro Ser Glu Ser Glu
530 535 540

Glu Glu Gln Ile Glu Ile Cys Ser Thr Arg Gly Cys Pro Asn Pro Pro
545 550 555 560

Ser Ser Pro Ala His Leu Pro Thr Gln Ala Ala Ile Cys Pro Ala Ser
565 570 575

Ala Lys Val Leu Ser Leu Lys Tyr Ser Thr Pro Arg Glu Ser Val Ala
580 585 590

Ser Pro Arg Glu Lys Ala Ala Cys Leu Pro Gly Ser Tyr Thr Ser Gly
595 600 605

Pro Asp Ser Ser Gln Pro Ser Ser Leu Leu Glu Met Ser Ser Gln Glu
610 615 620

His Glu Thr His Ala Asp Ile Ser Thr Ser Gln Asn His Arg Pro Ser
625 630 635 640

Cys Ala Glu Glu Thr Thr Glu Val Thr Ser Ala Ser Ser Ala Met Glu
645 650 655

Asn Ser Pro Leu Ser Lys Val Ala Arg His Phe His Ser Pro Pro Ile
660 665 670

Ile Leu Ser Ser Pro Asn Met Val Asn Gly Leu Glu His Asp Leu Leu
675 680 685

Asp Asp Glu Thr Leu Asn Gln Tyr Glu Thr Ser Ile Asn Ala Ala Ala
690 695 700

Ser Leu Ser Ser Phe Ser Val Asp Val Pro Lys Asn Gly Glu Ser Val
705 710 715 720

Leu Glu Asn Leu His Ile Ser Glu Ser Gln Asp Leu Asp Asp Leu Leu
725 730 735

Gln Lys Pro Lys Met Ile Ala Arg Arg Pro Ile Met Ala Trp Phe Lys
 740 745 750

 Glu Ile Asn Lys His Asn Gln Gly Thr His Leu Arg Ser Lys Thr Glu
 755 760 765

 Lys Glu Gln Pro Leu Met Pro Ala Arg Ser Pro Asp Ser Lys Ile Gln
 770 775 780

 Met Val Ser Ser Ser Gln Lys Lys Gly Val Thr Val Pro His Ser Pro
 785 790 795 800

 Pro Gln Pro Lys Thr Asn Leu Glu Asn Lys Asp Leu Ser Lys Lys Ser
 805 810 815

 Pro Ala Glu Met Leu Leu Thr Asn Gly Gln Lys Ala Lys Cys Gly Pro
 820 825 830

 Lys Leu Lys Arg Leu Ser Leu Lys Gly Lys Ala Lys Val Asn Ser Glu
 835 840 845

 Ala Pro Ala Ala Asn Ala Val Lys Ala Gly Gly Thr Asp His Arg Lys
 850 855 860

 Pro Leu Ile Ser Pro Gln Thr Ser His Lys Thr Leu Ser Lys Ala Val
 865 870 875 880

 Ser Gln Arg Leu His Val Ala Asp His Glu Asp Pro Asp Arg Asn Thr
 885 890 895

 Thr Ala Ala Pro Arg Ser Pro Gln Cys Val Leu Glu Ser Lys Pro Pro
 900 905 910

 Leu Ala Thr Ser Gly Pro Leu Lys Pro Ser Val Ser Asp Thr Ser Ile
 915 920 925

 Arg Thr Phe Val Ser Pro Leu Thr Ser Pro Lys Pro Val Pro Glu Gln
 930 935 940

 Gly Met Trp Ser Arg Phe His Met Ala Val Leu Ser Glu Pro Asp Arg
 945 950 955 960

 Gly Cys Pro Thr Thr Pro Lys Ser Pro Lys Cys Arg Ala Glu Gly Arg
 965 970 975

 Ala Pro Arg Ala Asp Ser Gly Pro Val Ser Pro Ala Ala Ser Arg Asn
 980 985 990

 Gly Met Ser Val Ala Gly Asn Arg Gln Ser Glu Pro Arg Leu Ala Ser
 995 1000 1005

 His Val Ala Ala Asp Thr Ala Gln Pro Arg Pro Thr Gly Glu Lys Gly
 1010 1015 1020

 Gly Asn Ile Met Ala Ser Asp Arg Leu Glu Arg Thr Asn Gln Leu Lys
 1025 1030 1035 1040

Ile Val Glu Ile Ser Ala Glu Ala Val Ser Glu Thr Val Cys Gly Asn
 1045 1050 1055

 Lys Pro Ala Glu Ser Asp Arg Arg Gly Gly Cys Leu Ala Gln Gly Asn
 1060 1065 1070

 Cys Gln Glu Lys Ser Glu Ile Arg Leu Tyr Arg Gln Val Ala Glu Ser
 1075 1080 1085

 Ser Thr Ser His Pro Ser Ser Leu Pro Ser His Ala Ser Gln Ala Glu
 1090 1095 1100

 Gln Glu Met Ser Arg Ser Phe Ser Met Ala Lys Leu Ala Ser Ser Ser
 1105 1110 1115 1120

 Ser Ser Leu Gln Thr Ala Ile Arg Lys Ala Glu Tyr Ser Gln Gly Lys
 1125 1130 1135

 Ser Ser Leu Met Ser Asp Ser Arg Gly Val Pro Arg Asn Ser Ile Pro
 1140 1145 1150

 Gly Gly Pro Ser Gly Glu Asp His Leu Tyr Phe Thr Pro Arg Pro Ala
 1155 1160 1165

 Thr Arg Thr Tyr Ser Met Pro Ala Gln Phe Ser Ser His Phe Gly Arg
 1170 1175 1180

 Glu Gly His Pro Pro His Ser Leu Gly Arg Ser Arg Asp Ser Gln Val
 1185 1190 1195 1200

 Pro Val Thr Ser Ser Val Val Pro Glu Ala Lys Ala Ser Arg Gly Gly
 1205 1210 1215

 Leu Pro Ser Leu Ala Asn Gly Gln Gly Ile Tyr Ser Val Lys Pro Leu
 1220 1225 1230

 Leu Asp Thr Ser Arg Asn Leu Pro Ala Thr Asp Glu Gly Asp Ile Ile
 1235 1240 1245

 Ser Val Gln Glu Thr Ser Cys Leu Val Thr Asp Lys Ile Lys Val Thr
 1250 1255 1260

 Arg Arg His Tyr Cys Tyr Glu Gln Asn Trp Pro His Glu Ser Thr Ser
 1265 1270 1275 1280

 Phe Phe Ser Val Lys Gln Arg Ile Lys Ser Phe Glu Asn Leu Ala Asn
 1285 1290 1295

 Ala Asp Arg Pro Val Ala Lys Ser Gly Ala Ser Pro Phe Leu Ser Val
 1300 1305 1310

 Ser Ser Lys Pro Pro Ile Gly Arg Arg Ser Ser Gly Ser Ile Val Ser
 1315 1320 1325

 Gly Ser Leu Gly His Pro Gly Asp Ala Ala Ala Arg Leu Leu Arg Arg
 1330 1335 1340

Ser Leu Ser Ser Cys Ser Glu Asn Gln Ser Glu Ala Gly Thr Leu Leu
 1345 1350 1355 1360

 Pro Gln Met Ala Lys Ser Pro Ser Ile Met Thr Leu Thr Ile Ser Arg
 1365 1370 1375

 Gln Asn Pro Pro Glu Thr Ser Ser Lys Gly Ser Asp Ser Glu Leu Lys
 1380 1385 1390

 Lys Ser Leu Gly Pro Leu Gly Ile Pro Thr Pro Thr Met Thr Leu Ala
 1395 1400 1405

 Ser Pro Val Lys Arg Asn Lys Ser Ser Val Arg His Thr Gln Pro Ser
 1410 1415 1420

 Pro Val Ser Arg Ser Lys Leu Gln Glu Leu Arg Ala Leu Ser Met Pro
 1425 1430 1435 1440

 Asp Leu Asp Lys Leu Cys Ser Glu Asp Tyr Ser Ala Gly Pro Ser Ala
 1445 1450 1455

 Val Leu Phe Lys Thr Glu Leu Glu Ile Thr Pro Arg Arg Ser Pro Gly
 1460 1465 1470

 Pro Pro Ala Gly Gly Val Ser Cys Pro Glu Lys Gly Asn Arg Ala
 1475 1480 1485

 Cys Pro Gly Gly Ser Gly Pro Lys Thr Ser Ala Ala Glu Thr Pro Ser
 1490 1495 1500

 Ser Ala Ser Asp Thr Gly Glu Ala Ala Gln Asp Leu Pro Phe Arg Arg
 1505 1510 1515 1520

 Ser Trp Ser Val Lys Leu Asp Gln Leu Leu Val Ser Ala Gly Asp Gln
 1525 1530 1535

 Gln Arg Leu Gln Ser Val Leu Ser Ser Val Gly Ser Lys Ser Thr Ile
 1540 1545 1550

 Leu Thr Leu Ile Gln Glu Ala Lys Ala Gln Ser Glu Asn Glu Glu Asp
 1555 1560 1565

 Val Cys Phe Ile Val Leu Asn Arg Lys Glu Gly Ser Gly Leu Gly Phe
 1570 1575 1580

 Ser Val Ala Gly Gly Thr Asp Val Glu Pro Lys Ser Ile Thr Val His
 1585 1590 1595 1600

 Arg Val Phe Ser Gln Gly Ala Ala Ser Gln Glu Gly Thr Met Asn Arg
 1605 1610 1615

 Gly Asp Phe Leu Leu Ser Val Asn Gly Ala Ser Leu Ala Gly Leu Ala
 1620 1625 1630

 His Gly Asn Val Leu Lys Val Leu His Gln Ala Gln Leu His Lys Asp
 1635 1640 1645

Ala Leu Val Val Ile Lys Lys Gly Met Asp Gln Pro Arg Pro Ser Ala
1650 1655 1660

Arg Gln Glu Pro Pro Thr Ala Asn Gly Lys Gly Leu Leu Ser Arg Lys
1665 1670 1675 1680

Thr Ile Pro Leu Glu Pro Gly Ile Gly Arg Ser Val Ala Val His Asp
1685 1690 1695

Ala Leu Cys Val Glu Val Leu Lys Thr Ser Ala Gly Leu Gly Leu Ser
1700 1705 1710

Leu Asp Gly Gly Lys Ser Ser Val Thr Gly Asp Gly Pro Leu Val Ile
1715 1720 1725

Lys Arg Val Tyr Lys Gly Gly Ala Ala Glu Gln Ala Gly Ile Ile Glu
1730 1735 1740

Ala Gly Asp Glu Ile Leu Ala Ile Asn Gly Lys Pro Leu Val Gly Leu
1745 1750 1755 1760

Met His Phe Asp Ala Trp Asn Ile Met Lys Ser Val Pro Glu Gly Pro
1765 1770 1775

Val Gln Leu Leu Ile Arg Lys His Arg Asn Ser Ser
1780 1785

<210> 61
<211> 1608
<212> PRT
<213> Homo sapiens

<400> 61
Ser Ser Asp Leu Ile Ser Ser Pro Gly Lys Lys Gly Ala Ala His Pro
1 5 10 15

Asp Pro Ser Lys Thr Ser Val Asp Thr Gly Lys Val Ser Arg Pro Glu
20 25 30

Asn Pro Ser Gln Pro Ala Ser Pro Arg Val Ala Lys Cys Lys Ala Arg
35 40 45

Ser Pro Val Arg Leu Pro His Glu Gly Ser Pro Ser Pro Gly Glu Lys
50 55 60

Ala Ala Ala Pro Pro Asp Tyr Ser Lys Thr Arg Ser Ala Ser Glu Thr
65 70 75 80

Ser Thr Pro His Asn Thr Arg Arg Val Ala Ala Leu Arg Gly Ala Gly
85 90 95

Pro Gly Ala Glu Gly Met Thr Pro Ala Gly Ala Val Leu Pro Gly Asp
100 105 110

Pro Leu Thr Ser Gln Glu Gln Arg Gln Gly Ala Pro Gly Asn His Ser
115 120 125

Lys Ala Leu Glu Met Thr Gly Ile His Ala Pro Glu Ser Ser Gln Glu
 130 135 140
 Pro Ser Leu Leu Glu Gly Ala Asp Ser Val Ser Ser Arg Ala Pro Gln
 145 150 155 160
 Ala Ser Leu Ser Met Leu Pro Ser Thr Asp Asn Thr Lys Glu Ala Cys
 165 170 175
 Gly His Val Ser Gly His Cys Cys Pro Gly Gly Ser Arg Glu Ser Pro
 180 185 190
 Val Thr Asp Ile Asp Ser Phe Ile Lys Glu Leu Asp Ala Ser Ala Ala
 195 200 205
 Arg Ser Pro Ser Ser Gln Thr Gly Asp Ser Gly Ser Gln Glu Gly Ser
 210 215 220
 Ala Gln Gly His Pro Pro Ala Gly Ala Gly Gly Ser Ser Cys Arg
 225 230 235 240
 Ala Glu Pro Val Pro Gly Gly Gln Thr Ser Ser Pro Arg Arg Ala Trp
 245 250 255
 Ala Ala Gly Ala Pro Ala Tyr Pro Gln Trp Ala Ser Gln Pro Ser Val
 260 265 270
 Leu Asp Ser Ile Asn Pro Asp Lys His Phe Thr Val Asn Lys Asn Phe
 275 280 285
 Leu Ser Asn Tyr Ser Arg Asn Phe Ser Ser Phe His Glu Asp Ser Thr
 290 295 300
 Ser Leu Ser Gly Leu Gly Asp Ser Thr Glu Pro Ser Leu Ser Ser Met
 305 310 315 320
 Tyr Gly Asp Ala Glu Asp Ser Ser Asp Pro Glu Ser Leu Thr Glu
 325 330 335
 Ala Pro Arg Ala Ser Ala Arg Asp Gly Trp Ser Pro Pro Arg Ser Arg
 340 345 350
 Val Ser Leu His Lys Glu Asp Pro Ser Glu Ser Glu Glu Gln Ile
 355 360 365
 Glu Ile Cys Ser Thr Arg Gly Cys Pro Asn Pro Pro Ser Ser Pro Ala
 370 375 380
 His Leu Pro Thr Gln Ala Ala Ile Cys Pro Ala Ser Ala Lys Val Leu
 385 390 395 400
 Ser Leu Lys Tyr Ser Thr Pro Arg Glu Ser Val Ala Ser Pro Arg Glu
 405 410 415
 Lys Val Ala Cys Leu Pro Gly Ser Tyr Thr Ser Gly Pro Asp Ser Ser
 420 425 430

Gln Pro Ser Ser Leu Leu Glu Met Ser Ser Gln Glu His Glu Thr His
 435 440 445
 Ala Asp Ile Ser Thr Ser Gln Asn His Arg Pro Ser Cys Ala Glu Glu
 450 455 460
 Thr Thr Glu Val Thr Ser Ala Ser Ser Ala Met Glu Asn Ser Pro Leu
 465 470 475 480
 Ser Lys Val Ala Arg His Phe His Ser Pro Pro Ile Ile Leu Ser Ser
 485 490 495
 Pro Asn Met Val Asn Gly Leu Glu His Asp Leu Leu Asp Asp Glu Thr
 500 505 510
 Leu Asn Gln Tyr Glu Thr Ser Ile Asn Ala Ala Ser Leu Ser Ser
 515 520 525
 Phe Ser Val Asp Val Pro Lys Asn Gly Glu Ser Val Leu Glu Asn Leu
 530 535 540
 His Ile Ser Glu Ser Gln Asp Leu Asp Asp Leu Leu Gln Lys Pro Lys
 545 550 555 560
 Met Ile Ala Arg Arg Pro Ile Met Ala Trp Phe Lys Glu Ile Asn Lys
 565 570 575
 His Asn Gln Gly Thr His Leu Arg Ser Lys Thr Glu Lys Glu Gln Pro
 580 585 590
 Leu Met Pro Ala Arg Ser Pro Asp Ser Lys Ile Gln Met Val Ser Ser
 595 600 605
 Ser Gln Lys Lys Gly Val Thr Val Pro His Ser Pro Pro Gln Pro Lys
 610 615 620
 Thr Asn Leu Glu Asn Lys Asp Leu Ser Lys Lys Ser Pro Ala Glu Met
 625 630 635 640
 Leu Leu Thr Asn Gly Gln Lys Ala Lys Cys Gly Pro Lys Leu Lys Arg
 645 650 655
 Leu Ser Leu Lys Gly Lys Ala Lys Val Asn Ser Glu Ala Pro Ala Ala
 660 665 670
 Asn Ala Val Lys Ala Gly Gly Thr Asp His Arg Lys Pro Leu Ile Ser
 675 680 685
 Pro Gln Thr Ser His Lys Thr Leu Ser Lys Ala Val Ser Gln Arg Leu
 690 695 700
 His Val Ala Asp His Glu Asp Pro Asp Arg Asn Thr Thr Ala Ala Pro
 705 710 715 720
 Arg Ser Pro Gln Cys Val Leu Glu Ser Lys Pro Pro Leu Ala Thr Ser
 725 730 735

Gly Pro Leu Lys Pro Ser Val Ser Asp Thr Ser Ile Arg Thr Phe Val
740 745 750

Ser Pro Leu Thr Ser Pro Lys Pro Val Pro Glu Gln Gly Met Trp Ser
755 760 765

Arg Phe His Met Ala Val Leu Ser Glu Pro Asp Arg Gly Cys Pro Thr
770 775 780

Thr Pro Lys Ser Pro Lys Cys Arg Ala Glu Gly Arg Ala Pro Arg Ala
785 790 795 800

Asp Ser Gly Pro Val Ser Pro Ala Ala Ser Arg Asn Gly Met Ser Val
805 810 815

Ala Gly Asn Arg Gln Ser Glu Pro Arg Leu Ala Ser His Val Ala Ala
820 825 830

Asp Thr Ala Gln Pro Arg Pro Thr Gly Glu Lys Gly Asn Ile Met
835 840 845

Ala Ser Asp Arg Leu Glu Arg Thr Asn Gln Leu Lys Ile Val Glu Ile
850 855 860

Ser Ala Glu Ala Val Ser Glu Thr Val Cys Gly Asn Lys Pro Ala Glu
865 870 875 880

Ser Asp Arg Arg Gly Gly Cys Leu Ala Gln Gly Asn Cys Gln Glu Lys
885 890 895

Ser Glu Ile Arg Leu Tyr Arg Gln Val Ala Glu Ser Ser Thr Ser His
900 905 910

Pro Ser Ser Leu Pro Ser His Ala Ser Gln Ala Glu Gln Glu Met Ser
915 920 925

Arg Ser Phe Ser Met Ala Lys Leu Ala Ser Ser Ser Ser Ser Leu Gln
930 935 940

Thr Ala Ile Arg Lys Ala Glu Tyr Ser Gln Gly Lys Ser Ser Leu Met
945 950 955 960

Ser Asp Ser Arg Gly Val Pro Arg Asn Ser Ile Pro Gly Gly Pro Ser
965 970 975

Gly Glu Asp His Leu Tyr Phe Thr Pro Arg Pro Ala Thr Arg Thr Tyr
980 985 990

Ser Met Pro Ala Gln Phe Ser Ser His Phe Gly Arg Glu Gly His Pro
995 1000 1005

Pro His Ser Leu Gly Arg Ser Arg Asp Ser Gln Val Pro Val Thr Ser
1010 1015 1020

Ser Val Val Pro Glu Ala Lys Ala Ser Arg Gly Gly Leu Pro Ser Leu
1025 1030 1035 1040

Ala Asn Gly Gln Gly Ile Tyr Ser Val Lys Pro Leu Leu Asp Thr Ser
1045 1050 1055

Arg Asn Leu Pro Ala Thr Asp Glu Gly Asp Ile Ile Ser Val Gln Glu
1060 1065 1070

Thr Ser Cys Leu Val Thr Asp Lys Ile Lys Val Thr Arg Arg His Tyr
1075 1080 1085

Cys Tyr Glu Gln Asn Trp Pro His Glu Ser Thr Ser Phe Phe Ser Val
1090 1095 1100

Lys Gln Arg Ile Lys Ser Phe Glu Asn Leu Ala Asn Ala Asp Arg Pro
1105 1110 1115 1120

Val Ala Lys Ser Gly Ala Ser Pro Phe Leu Ser Val Ser Ser Lys Pro
1125 1130 1135

Pro Ile Gly Arg Arg Ser Ser Gly Ser Ile Val Ser Gly Ser Leu Gly
1140 1145 1150

His Pro Gly Asp Ala Ala Ala Arg Leu Leu Arg Arg Ser Leu Ser Ser
1155 1160 1165

Cys Ser Glu Asn Gln Ser Glu Ala Gly Thr Leu Leu Pro Gln Met Ala
1170 1175 1180

Lys Ser Pro Ser Ile Met Thr Leu Thr Ile Ser Arg Gln Asn Pro Pro
1185 1190 1195 1200

Glu Thr Ser Ser Lys Gly Ser Asp Ser Glu Leu Lys Lys Ser Leu Gly
1205 1210 1215

Pro Leu Gly Ile Pro Thr Pro Thr Met Thr Leu Ala Ser Pro Val Lys
1220 1225 1230

Arg Asn Lys Ser Ser Val Arg His Thr Gln Pro Ser Pro Val Ser Arg
1235 1240 1245

Ser Lys Leu Gln Glu Leu Arg Ala Leu Ser Met Pro Asp Leu Asp Lys
1250 1255 1260

Leu Cys Ser Glu Asp Tyr Ser Ala Gly Pro Ser Ala Val Leu Phe Lys
1265 1270 1275 1280

Thr Glu Leu Glu Ile Thr Pro Arg Arg Ser Pro Gly Pro Pro Ala Gly
1285 1290 1295

Gly Val Ser Cys Pro Glu Lys Gly Gly Asn Arg Ala Cys Pro Gly Gly
1300 1305 1310

Ser Gly Pro Lys Thr Ser Ala Ala Glu Thr Pro Ser Ser Ala Ser Asp
1315 1320 1325

Thr Gly Glu Ala Ala Gln Asp Leu Pro Phe Arg Arg Ser Trp Ser Val
1330 1335 1340

Asn Leu Asp Gln Leu Leu Val Ser Ala Gly Asp Gln Gln Arg Leu Gln
 1345 1350 1355 1360
 Ser Val Leu Ser Ser Val Gly Ser Lys Ser Thr Ile Leu Thr Leu Ile
 1365 1370 1375
 Gln Glu Ala Lys Ala Gln Ser Glu Asn Glu Glu Asp Val Cys Phe Ile
 1380 1385 1390
 Val Leu Asn Arg Lys Glu Gly Ser Gly Leu Gly Phe Ser Val Ala Gly
 1395 1400 1405
 Gly Thr Asp Val Glu Pro Lys Ser Ile Thr Val His Arg Val Phe Ser
 1410 1415 1420
 Gln Gly Ala Ala Ser Gln Glu Gly Thr Met Asn Arg Gly Asp Phe Leu
 1425 1430 1435 1440
 Leu Ser Val Asn Gly Ala Ser Leu Ala Gly Leu Ala His Gly Asn Val
 1445 1450 1455
 Leu Lys Val Leu His Gln Ala Gln Leu His Lys Asp Ala Leu Val Val
 1460 1465 1470
 Ile Lys Lys Gly Met Asp Gln Pro Arg Pro Ser Ala Arg Gln Glu Pro
 1475 1480 1485
 Pro Thr Ala Asn Gly Lys Gly Leu Leu Ser Arg Lys Thr Ile Pro Leu
 1490 1495 1500
 Glu Pro Gly Ile Gly Arg Ser Val Ala Val His Asp Ala Leu Cys Val
 1505 1510 1515 1520
 Glu Val Leu Lys Thr Ser Ala Gly Leu Gly Leu Ser Leu Asp Gly Gly
 1525 1530 1535
 Lys Ser Ser Val Thr Gly Asp Gly Pro Leu Val Ile Lys Arg Val Tyr
 1540 1545 1550
 Lys Gly Gly Ala Ala Glu Gln Ala Gly Ile Ile Glu Ala Gly Asp Glu
 1555 1560 1565
 Ile Leu Ala Ile Asn Gly Lys Pro Leu Val Gly Leu Met His Phe Asp
 1570 1575 1580
 Ala Trp Asn Ile Met Lys Ser Val Pro Glu Gly Pro Val Gln Leu Leu
 1585 1590 1595 1600
 Ile Arg Lys His Arg Asn Ser Ser
 1605

<210> 62
 <211> 2766
 <212> PRT
 <213> Rattus norvegicus

<400> 62
Met Pro Ile Thr Gln Asp Asn Ala Leu Leu His Leu Pro Leu Leu Tyr
1 5 10 15

Glu Trp Leu Gln Asn Ser Leu Arg Glu Gly Gly Asp Ser Pro Glu Gln
20 25 30

Arg Leu Cys Gln Ala Ala Ile Gln Lys Leu Gln Glu Tyr Ile Gln Leu
35 40 45

Asn Leu Ala Val Asp Glu Ser Thr Val Pro Pro Asp His Ser Pro Pro
50 55 60

Glu Met Glu Ile Cys Thr Val Tyr Leu Thr Lys Gln Leu Gly Asp Thr
65 70 75 80

Glu Thr Val Gly Leu Ser Phe Gly Asn Ile Pro Val Phe Gly Asp Tyr
85 90 95

Gly Glu Lys Arg Arg Gly Gly Lys Lys Arg Lys Thr His Gln Gly Pro
100 105 110

Val Leu Asp Val Gly Cys Ile Trp Val Thr Glu Leu Arg Lys Asn Ser
115 120 125

Pro Ala Gly Lys Ser Gly Lys Val Arg Leu Arg Asp Glu Ile Leu Ser
130 135 140

Leu Asn Gly Gln Leu Met Val Gly Val Asp Val Thr Gly Ala Ser Tyr
145 150 155 160

Leu Ala Glu Gln Cys Trp Asn Gly Phe Ile Tyr Leu Ile Met Leu
165 170 175

Arg Arg Phe Lys Gln Lys Ala His Val Thr Tyr Asn Gly Asn Ser Gly
180 185 190

Asn Ser Ser Glu Pro Gly Glu Thr Pro Thr Leu Glu Leu Gly Asp Gln
195 200 205

Thr Ser Lys Lys Gly Lys Arg Thr Arg Lys Phe Gly Val Ile Ser Arg
210 215 220

Pro Ser Ile Ser Lys Thr Pro Glu Asp Ser Lys Ser Ser Ser Gly Cys
225 230 235 240

Asp Thr Ala Asp Asp Pro Asn Ser Glu Leu Glu Asn Gly Ala Asp Pro
245 250 255

Glu Leu Gly Asn Gly His Ala Phe Glu Leu Glu Asn Gly Pro His Ser
260 265 270

Leu Lys Asp Val Ala Gly Pro His Leu Glu Arg Ser Glu Ala Asp Ser
275 280 285

Glu Val Glu Leu Arg Val Pro Lys Thr Glu Ala Pro Leu Ser Asp Ser

290

295

300

Asn Asp Lys Arg Arg Phe Ser Lys Thr Gly Lys Thr Asp Phe Gln Ser
 305 310 315 320

Ser Asp Cys Leu Ala Arg Glu Glu Val Gly Arg Ile Trp Lys Met Glu
 325 330 335

Leu Leu Lys Glu Ser Asp Gly Leu Gly Ile Gln Val Ser Gly Gly Arg
 ' 340 345 350

Gly Ser Lys Arg Ser Pro His Ala Ile Val Val Thr Gln Val Lys Glu
 355 360 365

Gly Gly Ala Ala His Arg Asp Gly Arg Leu Ser Leu Gly Asp Glu Leu
 370 375 380

Leu Val Ile Asn Gly His Leu Leu Val Gly Leu Ser His Glu Glu Ala
 385 390 395 400

Val Ala Ile Leu Arg Ser Ala Thr Gly Met Val Gln Leu Val Val Ala
 405 410 415

Ser Lys Met Pro Gly Ser Glu Glu Ser Gln Asp Val Gly Ser Ser Glu
 420 425 430

Glu Ser Lys Gly Asn Leu Glu Ser Pro Lys Gln Gly Asn Cys Lys Thr
 435 440 445

Lys Leu Lys Ser Arg Leu Ser Gly Gly Val His Arg Leu Glu Ser Val
 450 455 460

Glu Glu Tyr Asn Glu Leu Met Val Arg Asn Gly Asp Pro Arg Ile Arg
 465 470 475 480

Met Leu Glu Val Ser Arg Asp Gly Arg Lys His Ser Leu Pro Gln Leu
 485 490 495

Leu Asp Ser Thr Gly Thr Ser Gln Glu Tyr His Ile Val Lys Lys Ser
 500 505 510

Thr Arg Ser Leu Ser Thr Thr His Val Glu Ser Pro Trp Arg Leu Ile
 515 520 525

Arg Pro Ser Val Ile Ser Ile Ile Gly Leu Tyr Lys Glu Lys Gly Lys
 530 535 540

Gly Leu Gly Phe Ser Ile Ala Gly Gly Arg Asp Cys Ile Arg Gly Gln
 545 550 555 560

Met Gly Ile Phe Val Lys Thr Ile Phe Pro Asn Gly Ser Ala Ala Glu
 565 570 575

Asp Gly Arg Leu Lys Glu Gly Asp Glu Ile Leu Asp Val Asn Gly Ile
 580 585 590

Pro Ile Lys Gly Leu Thr Phe Gln Glu Ala Ile His Thr Phe Lys Gln

595	600	605
Ile Arg Ser Gly Leu Phe Val Leu Thr Val Arg Thr Lys Leu Leu Ser		
610	615	620
Pro Ser Leu Thr Pro Cys Ser Thr Pro Thr His Met Ser Arg Ser Ser		
625	630	635
Ser Pro Ser Phe Asn Thr Asn Ser Gly Gly Thr Pro Ala Gly Gly Gly		
645	650	655
Gln Glu Glu Gly Gly Ser Ser Ser Leu Gly Arg Lys Ala Pro Gly Pro		
660	665	670
Lys Asp Arg Ile Val Met Glu Val Thr Leu Asn Lys Glu Pro Arg Val		
675	680	685
Gly Leu Gly Ile Gly Ala Cys Cys Leu Ala Leu Glu Asn Ser Pro Pro		
690	695	700
Gly Ile Tyr Ile His Ser Leu Ala Pro Gly Ser Val Ala Lys Met Glu		
705	710	715
Ser Asn Leu Ser Arg Gly Asp Gln Ile Leu Glu Val Asn Ser Val Asn		
725	730	735
Val Arg His Ala Ala Leu Ser Lys Val His Ala Ile Leu Ser Lys Cys		
740	745	750
Pro Pro Gly Pro Val Arg Leu Val Ile Gly Arg His Pro Asn Pro Lys		
755	760	765
Val Ser Glu Gln Glu Met Asp Glu Val Ile Ala Arg Ser Thr Tyr Gln		
770	775	780
Glu Ser Arg Glu Ala Asn Ser Ser Pro Gly Leu Gly Thr Pro Leu Lys		
785	790	795
Ser Pro Ser Leu Ala Lys Lys Asp Ser Leu Leu Ser Glu Ser Glu Leu		
805	810	815
Ser Gln Tyr Phe Val His Asp Gly Gln Gly Ser Leu Ser Asp Phe Val		
820	825	830
Val Ala Gly Ser Glu Asp Glu Asp His Pro Gly Ser Gly Tyr Glu Thr		
835	840	845
Ser Glu Asp Gly Ser Leu Leu Pro Val Pro Ser Ala His Lys Ala Arg		
850	855	860
Ala Asn Ser Leu Val Thr Leu Gly Ser Gln Arg Thr Ser Gly Leu Leu		
865	870	875
His Lys Gln Val Thr Val Ala Arg Gln Ala Ser Leu Pro Gly Ser Pro		
885	890	895
Gln Val Leu Arg Asn Pro Leu Leu Arg Gln Arg Arg Val Arg Cys Tyr		

900	905	910
Asp Ser Asn Gly Gly Ser Asp Asp Glu Asp Phe Asp Gly Glu Gly Asp		
915	920	925
Cys Ile Ser Leu Pro Gly Val Leu Pro Gly Pro Gly Lys Pro Leu Val		
930	935	940
Glu Asp Asp Thr Arg Pro Ala Leu Thr Thr Ser Ser Lys Ser Ile Asp		
945	950	955
Val Asn Lys Gln Glu Glu Arg Leu Gln Lys Pro Leu Val Ser Lys Ala		
965	970	975
Cys Ser Val Pro Leu Leu Gly Ser Ser Leu Asp Ser Glu His Ser Ile		
980	985	990
Leu Asn Gly Ala Gly Gly Thr Pro Pro Lys Val Ala Ser Leu Pro Gly		
995	1000	1005
Ser Gly Glu Thr Pro Lys Asn Gly Pro Arg Gly Ser Gly Arg Lys Glu		
1010	1015	1020
Met Ser Gly Ser Arg Ser Ser Pro Lys Leu Glu Tyr Arg Val Pro Thr		
1025	1030	1035
Asp Thr Gln Ser Pro Arg Ser Pro Glu Asn His Thr Ser Pro Pro Gln		
1045	1050	1055
Lys Ser Glu Asn Leu Val Ser Arg His Lys Pro Val Ala Arg Ile Ser		
1060	1065	1070
Pro His Tyr Lys Arg Ser Asp Ala Glu Glu Ala Pro Gly Gly Thr Ala		
1075	1080	1085
Asn Gly Pro Cys Ala Gln Asp Leu Lys Val Gln Ala Ser Pro Val Lys		
1090	1095	1100
Asp Pro Val Thr Ser Arg Gln Pro Gly Gly Thr Ala Glu Lys Glu Leu		
1105	1110	1115
1120		
Arg Gly Asn Pro Thr Pro Gly Asp Ser Ser Val Pro Thr Asn Cys Gly		
1125	1130	1135
Pro Ala Ser Thr Pro Cys His Pro Asn Ile Gly Leu Pro Thr Glu Asn		
1140	1145	1150
Pro Gln Gly Ala Ala Pro Glu Cys Gly Pro His Pro Gly Thr Gly Trp		
1155	1160	1165
Asp Gly Ser Ser Glu His Leu Cys Ser Pro Gly Lys Ser Arg Glu Val		
1170	1175	1180
His Pro Asp Ser Ser Glu Thr Pro Thr Val Ala Glu Gln Val His Gln		
1185	1190	1195
1200		
Pro Glu Ser Leu Ser Gln Pro Val Ser Pro Arg Thr Ser Glu Pro Glu		

1205	1210	1215	
Ser Gln Gly Ile Ser Lys Met Lys Pro Pro Ser Gln Arg Cys Val Ser			
1220	1225	1230	
Pro Arg Glu Lys Ala Ser Thr Pro Pro Asp Ser Ser Arg Ala Trp Ala			
1235	1240	1245	
Ala Pro Gly Asp Ser Ser Pro Ser Thr Arg Arg Ile Ala Val Pro Met			
1250	1255	1260	
Ser Thr Gly Ala Ala Pro Ala Thr Ala Ile Pro Gln Ala Ser Leu Val			
1265	1270	1275	1280
Ser Gln Glu Arg Ser Arg Gly Leu Ser Gly Pro Ser Lys Gly Leu Gly			
1285	1290	1295	
Thr Lys Glu Leu Cys Ile Pro Lys Ser Leu Lys Asp Gly Ala Leu Leu			
1300	1305	1310	
Glu Asp Thr Ala Pro Ala Ser Gly Lys Met Ser His Ala Ser Ser Pro			
1315	1320	1325	
Ser Gly Pro Val Ala Thr Glu Arg Thr Leu Ser Gly Ser Pro Glu Asn			
1330	1335	1340	
Pro Val Thr Asp Ile Asp Asn Phe Ile Glu Glu Ala Ser Glu Ala Arg			
1345	1350	1355	1360
Leu Ser Gln Ser Pro Gln Lys Ala Asp Cys Arg Ala His Gly Asp Thr			
1365	1370	1375	
Phe Glu Ser Gln Pro Pro Gly Gly Ala Gly Ser Ser Ser His His			
1380	1385	1390	
Ala Gln Met Val Arg Ser Asp Gln Thr Ser Ser Pro Arg Lys Thr Gly			
1395	1400	1405	
Gly Thr Gly Ser Pro Pro Gln Gln Trp Ala Leu Gln Pro Ser Val			
1410	1415	1420	
Leu Asp Ser Ile His Pro Asp Lys His Leu Ala Val Asn Lys Thr Phe			
1425	1430	1435	1440
Leu Asn Asn Tyr Ser Arg Asn Phe Ser Asn Phe His Glu Asp Ser Ile			
1445	1450	1455	
Ser Leu Ser Gly Pro Gly Gly Ser Ser Glu Pro Ser Pro Ser Ser Met			
1460	1465	1470	
Tyr Gly Asn Ala Glu Asp Ser Ser Asp Pro Glu Ser Leu Ala Glu			
1475	1480	1485	
Asp Pro Gly Ala Ala Ala Arg Asn Asn Trp Ser Pro Pro Leu Ser Pro			
1490	1495	1500	
Glu Ser Ser Pro Lys Glu Gly Ser Ser Glu Ser Glu Asp Glu Arg Ile			

1505	1510	1515	1520
Glu Ile Cys Ser Thr Asp Gly Cys Pro Gly Thr Pro Val Thr Ala Pro			
1525	1530	1535	
Pro Pro Thr Gln Val Ala Leu Cys Pro Val Leu Pro Val Gln Gln Arg			
1540	1545	1550	
Ala Val Cys Lys Pro Val Gly Asp Ile Cys Glu Arg Ala Cys Phe Val			
1555	1560	1565	
Pro Gly Ala Ser Arg Thr Ser Ile Pro Asp Ser Ser Gln Pro Phe Ser			
1570	1575	1580	
Phe Leu Asp Val Ser Ser Glu Glu Pro Glu Thr Trp Ala Ser Ile Asn			
1585	1590	1595	1600
Ala Ser Gln Asn His Met Pro Val Cys Thr Glu Gly Ile Met Asp Val			
1605	1610	1615	
Thr Ser Thr Ser Ser Asn Met Gly Asp Ser Gln Ser Ser Gln Met Thr			
1620	1625	1630	
Arg His Cys Arg Asn Ala Pro Phe Val Leu Gly Asn Pro Asp Met Val			
1635	1640	1645	
Asn Asp Leu Gly Arg Asp Leu Leu Asp Glu Gly Ala Pro Lys Glu Gly			
1650	1655	1660	
Ala Ala Ala Ala Ser Val Met Arg Ser Val Phe Ala Leu Gly Ala Glu			
1665	1670	1675	1680
Gly Pro Lys Asn Gly Glu Ala Val Leu Ala Asp Leu His Ile Ala Glu			
1685	1690	1695	
Arg Gly Asn Leu Glu Asp Leu Leu Gln Lys Pro Lys Thr Ile Ser Arg			
1700	1705	1710	
Arg Pro Ile Leu Thr Trp Phe Lys Glu Ile Asn Lys Asp Ser Gln Gly			
1715	1720	1725	
Ser His Leu Arg Ser Thr Ser Glu Lys Glu Gln Ser Ser Met Leu Ala			
1730	1735	1740	
Leu Gly Pro Gly Ser Lys Ala Asn Met Val Asn Thr Gly His Arg Lys			
1745	1750	1755	1760
Gly Val Thr Val Pro Lys Ser Pro Pro Ser Arg Gln Lys Ser Gln Glu			
1765	1770	1775	
Asn Lys Asp Leu Pro Pro Lys Ser Pro Val Glu Thr Leu Gly Asn Cys			
1780	1785	1790	
Gln Lys Pro Lys Cys Ser Pro Lys Leu Lys Arg Leu Asn Ser Lys Gly			
1795	1800	1805	
Lys Ala Ser Pro Glu Val Pro Val Ala Ile Ser Thr Lys Gly Ser Arg			

1810	1815	1820
Asn Asp His Arg Lys Thr Leu Pro Ser Pro Gln Ala Ser His Lys Met		
1825	1830	1835
Phe Ser Lys Ala Val Ser His Arg Leu His Ile Ala Asp Gln Glu Glu		
1845	1850	1855
Pro Lys Asn Thr Ala Gly Asp Thr Pro Lys Pro Pro Gln Cys Val Pro		
1860	1865	1870
Glu Ser Lys Pro Pro Gln Ala Ala Leu Gly Ser Leu Arg Thr Ser Ala		
1875	1880	1885
Ser Asp Thr Ser Ile Arg Thr Phe Thr Ser Pro Leu Thr Ser Pro Lys		
1890	1895	1900
Leu Leu Pro Glu Gln Gly Ala Asn Ser Arg Phe His Met Ala Val Tyr		
1905	1910	1915
Leu Glu Ser Asp Thr Ser Cys Pro Thr Thr Ser Arg Ser Pro Arg Ser		
1925	1930	1935
Gly Pro Glu Gly Lys Ala Pro His Ala Asn Ser Gly Ser Ala Ser Pro		
1940	1945	1950
Pro Ala Ser Arg Ala Ser Leu Ala Leu Ala Gly Ile Arg Gln Ser Lys		
1955	1960	1965
Gln Phe Thr Pro Gly Arg Ala Asp Leu Leu Val Ser Glu Ala Thr Gln		
1970	1975	1980
Pro Gln Gly Ile Cys Glu Lys Gly Ala Glu Lys Lys Val Ser Asp Pro		
1985	1990	1995
2000		
Pro Gln Arg Thr Asn Gln Leu Lys Ile Val Glu Ile Ser Ser Glu Arg		
2005	2010	2015
Val Pro Lys Asn Ala Cys Gly Asp Arg Pro Pro Glu Ser Asp Arg Lys		
2020	2025	2030
Gly Gly Phe Leu Thr Gln Asn Asn Cys Gln Glu Lys Ser Ala Ile Arg		
2035	2040	2045
Leu Arg Gln Ser Glu Glu Ser Ser Pro Glu His Thr Pro Phe Pro Pro		
2050	2055	2060
Ser Gln Ala Ser Gln Val Glu Arg Glu Ile Arg Trp Ser Phe Ser Met		
2065	2070	2075
2080		
Ala Lys Pro Ala Thr Ser Ser Ser Ser Leu Gln Leu Pro Ala Lys		
2085	2090	2095
Leu Pro Glu Ser Phe Gln Gly Lys Ser Ser Gln Met Pro Ala Ser Val		
2100	2105	2110
Gly Val Pro Lys Asn Gly Val Pro Ile Gly Leu Ala Gly Glu Glu Ser		

2115

2120

2125

Pro Tyr Phe Thr Pro Arg Pro Ala Thr Arg Thr Tyr Ser Met Pro Ala
 2130 2135 2140
 Gln Phe Ser Ser His Phe Gly Arg Glu Gly Pro Ser Pro His Ser Pro
 2145 2150 2155 2160
 Ser His Ser Pro Gln Asp Pro Gln Val Pro Ala Met Gly Gly Lys Leu
 2165 2170 2175
 Ser Glu Lys Thr Ala Lys Gly Val Thr Asn Gly Gln Gly Val Tyr Ser
 2180 2185 2190
 Val Lys Pro Leu Leu Glu Thr Ser Lys Asn Leu Ser Pro Val Asp Gly
 2195 2200 2205
 Arg Asp Val Ser Ala Asp Pro Glu Thr Ser Cys Leu Ile Pro Asp Lys
 2210 2215 2220
 Val Lys Val Thr Arg Arg Gln Tyr Cys Cys Glu Gln Ser Trp Pro His
 2225 2230 2235 2240
 Glu Ser Thr Ser Phe Phe Ser Val Lys Gln Arg Ile Lys Ser Phe Glu
 2245 2250 2255
 Asn Leu Ala Asn Ser Asp Arg Pro Thr Ala Lys Cys Ala Thr Ser Pro
 2260 2265 2270
 Phe Leu Ser Val Ser Ser Lys Pro Pro Ile Asn Arg Arg Ser Ser Gly
 2275 2280 2285
 Ser Ile Pro Ser Gly Ser Pro Ser Asp Met Thr Ser Arg Ser Leu Arg
 2290 2295 2300
 Arg Ser Leu Ser Ser Cys Ser Glu Ser Gln Ser Glu Ala Ser Ser Leu
 2305 2310 2315 2320
 Leu Pro Gln Met Thr Lys Ser Pro Ser Ser Met Thr Leu Thr Val Ser
 2325 2330 2335
 Arg Gln Asn Pro Pro Asp Thr Ser Asn Lys Gly Pro Ser Pro Asp Pro
 2340 2345 2350
 Lys Lys Ser Leu Val Pro Val Gly Ile Pro Thr Ser Thr Val Ser Pro
 2355 2360 2365
 Ala Ser Pro Ser Lys Arg Asn Lys Ser Ser Val Arg His Ala Gln Pro
 2370 2375 2380
 Ser Pro Val Ser Arg Ser Lys Leu Gln Glu Arg Arg Thr Leu Ser Met
 2385 2390 2395 2400
 Pro Asp Leu Asp Lys Leu Cys Asn Gly Glu Asp Asp Ser Ala Ser Pro
 2405 2410 2415
 Gly Ala Val Leu Phe Lys Thr Gln Leu Glu Ile Thr Pro Arg Arg Ser

	2420	2425	2430
Lys Gly Ser Gln Ala Thr Ser Pro Ala Gly Ser Pro Ala Arg Gly His			
2435	2440	2445	
Ala Asp Phe Asn Gly Ser Thr Phe Leu Ser Cys Pro Met Asn Gly Gly			
2450	2455	2460	
Thr Arg Ala Tyr Thr Lys Gly Asn Ser Pro Pro Ala Ser Glu Pro Ala			
2465	2470	2475	2480
Ile Ala Thr Gly Ser Arg Glu Glu Gly Ser Val Trp Ala Thr Pro			
2485	2490	2495	
Ser Gly Lys Ser Trp Ser Val Ser Leu Asp Arg Leu Leu Ala Ser Val			
2500	2505	2510	
Gly Asn Gln Gln Arg Leu Gln Gly Ile Leu Ser Leu Val Gly Ser Lys			
2515	2520	2525	
Ser Pro Ile Leu Thr Leu Ile Gln Glu Ala Lys Ala Gln Ser Glu Thr			
2530	2535	2540	
Lys Glu Asp Ile Cys Phe Ile Val Leu Asn Lys Lys Glu Gly Ser Gly			
2545	2550	2555	2560
Leu Gly Phe Ser Val Ala Gly Gly Ala Asp Val Glu Pro Lys Ser Val			
2565	2570	2575	
Met Val His Arg Val Phe Ser Gln Gly Val Ala Ser Gln Glu Gly Thr			
2580	2585	2590	
Val Ser Arg Gly Asp Phe Leu Leu Ser Val Asn Gly Thr Ser Leu Ala			
2595	2600	2605	
Gly Leu Ala His Ser Glu Val Thr Lys Val Leu His Gln Ala Glu Leu			
2610	2615	2620	
His Lys His Ala Leu Met Ile Ile Lys Lys Gly Asn Asp Gln Pro Gly			
2625	2630	2635	2640
Pro Ser Phe Lys Gln Glu Pro Pro Ser Ala Asn Gly Lys Gly Pro Phe			
2645	2650	2655	
Pro Arg Arg Thr Leu Pro Leu Glu Pro Gly Ala Gly Arg Asn Gly Ala			
2660	2665	2670	
Ala His Asp Ala Leu Cys Val Glu Val Leu Lys Thr Ser Ala Gly Leu			
2675	2680	2685	
Gly Leu Ser Leu Asp Gly Gly Lys Ser Ser Val Ser Gly Glu Gly Pro			
2690	2695	2700	
Leu Val Ile Lys Arg Val Tyr Lys Gly Gly Ala Ala Glu Arg Ala Gly			
2705	2710	2715	2720
Thr Ile Glu Ala Gly Asp Glu Ile Leu Ala Ile Asn Gly Lys Pro Leu			

2725

2730

2735

Val Gly Leu Val His Phe Asp Ala Trp Asn Ile Met Lys Ser Val Pro
 2740 2745 2750

Glu Gly Pro Val Gln Leu Val Ile Arg Lys His Arg Asp Ser
 2755 2760 2765

<210> 63
<211> 2641
<212> PRT
<213> Homo sapiens

<400> 63
Met Pro Ile Thr Gln Asp Asn Ala Val Leu His Leu Pro Leu Leu Tyr
 1 5 10 15

Gln Trp Leu Gln Asn Ser Leu Gln Glu Gly Gly Asp Gly Pro Glu Gln
 20 25 30

Arg Leu Cys Gln Ala Ala Ile Gln Lys Leu Gln Glu Tyr Ile Gln Leu
 35 40 45

Asn Phe Ala Val Asp Glu Ser Thr Val Pro Pro Asp His Ser Pro Pro
 50 55 60

Glu Met Glu Ile Cys Thr Val Tyr Leu Thr Lys Glu Leu Gly Asp Thr
 65 70 75 80

Glu Thr Val Gly Leu Ser Phe Gly Asn Ile Pro Val Phe Gly Asp Tyr
 85 90 95

Gly Glu Lys Arg Arg Gly Gly Lys Arg Lys Thr His Gln Gly Pro
 100 105 110

Val Leu Asp Val Gly Cys Ile Trp Val Thr Glu Leu Arg Lys Asn Ser
 115 120 125

Pro Ala Gly Lys Ser Gly Lys Val Arg Leu Arg Asp Glu Ile Leu Ser
 130 135 140

Leu Asn Gly Gln Leu Met Val Gly Val Asp Val Ser Gly Ala Ser Tyr
 145 150 155 160

Leu Ala Glu Gln Cys Trp Asn Gly Gly Phe Ile Tyr Leu Ile Met Leu
 165 170 175

Arg Arg Phe Lys His Lys Ala His Ser Thr Tyr Asn Gly Asn Ser Ser
 180 185 190

Asn Ser Ser Glu Pro Gly Glu Thr Pro Thr Leu Glu Leu Gly Asp Arg
 195 200 205

Thr Ala Lys Lys Gly Lys Arg Thr Arg Lys Phe Gly Val Ile Ser Arg
 210 215 220

Pro Pro Ala Asn Lys Ala Pro Glu Glu Ser Lys Gly Ser Ala Gly Cys
225 230 235 240

Glu Val Ser Ser Asp Pro Ser Thr Glu Leu Glu Asn Gly Ala Asp Pro
245 250 255

Glu Leu Gly Asn Gly His Val Phe Gln Leu Glu Asn Gly Pro Asp Ser
260 265 270

Leu Lys Glu Val Ala Gly Pro His Leu Glu Arg Ser Glu Val Asp Arg
275 280 285

Gly Thr Glu His Arg Ile Pro Lys Thr Asp Ala Pro Leu Thr Thr Ser
290 295 300

Asn Asp Lys Arg Arg Phe Ser Lys Gly Gly Lys Thr Asp Phe Gln Ser
305 310 315 320

Ser Asp Cys Leu Ala Arg Gln Glu Glu Val Gly Arg Ile Trp Lys Met
325 330 335

Glu Leu Leu Lys Glu Ser Asp Gly Leu Gly Ile Gln Val Ser Gly Gly
340 345 350

Arg Gly Ser Lys Arg Ser Pro His Ala Ile Val Val Thr Gln Val Lys
355 360 365

Glu Gly Gly Ala Ala His Arg Glu Tyr His Ile Val Lys Ser Thr
370 375 380

Arg Ser Leu Ser Thr Thr Gln Val Glu Ser Pro Trp Arg Leu Ile Arg
385 390 395 400

Pro Ser Val Ile Ser Ile Ile Gly Leu Tyr Lys Glu Lys Gly Lys Gly
405 410 415

Leu Gly Phe Ser Ile Ala Gly Gly Arg Asp Cys Ile Arg Gly Gln Met
420 425 430

Gly Ile Phe Val Lys Thr Ile Phe Pro Asn Gly Ser Ala Ala Glu Asp
435 440 445

Gly Arg Leu Lys Glu Gly Asp Glu Ile Leu Asp Val Asn Gly Ile Pro
450 455 460

Ile Lys Gly Leu Thr Phe Gln Glu Ala Ile His Thr Phe Lys Gln Ile
465 470 475 480

Arg Ser Gly Leu Phe Val Leu Thr Val Arg Thr Lys Leu Val Ser Pro
485 490 495

Ser Leu Thr Pro Cys Ser Thr Pro Thr His Met Ser Arg Ser Ala Ser
500 505 510

Pro Asn Phe Asn Thr Ser Gly Gly Ala Ser Ala Gly Ser Asp Glu
515 520 525

Gly Ser Ser Ser Leu Gly Arg Lys Thr Pro Gly Pro Lys Asp Arg
 530 535 540
 Ile Val Met Glu Val Thr Leu Asn Lys Glu Pro Arg Val Gly Leu Gly
 545 550 555 560
 Ile Gly Ala Cys Cys Leu Ala Leu Glu Asn Ser Pro Pro Gly Ile Tyr
 565 570 575
 Ile His Ser Leu Ala Pro Gly Ser Val Ala Lys Met Glu Ser Asn Leu
 580 585 590
 Ser Arg Gly Asp Gln Ile Leu Glu Val Asn Ser Val Asn Val Arg His
 595 600 605
 Ala Ala Leu Ser Lys Val His Ala Ile Leu Ser Lys Cys Pro Val Ser
 610 615 620
 Glu Gln Glu Met Asp Glu Val Ile Ala Arg Ser Thr Tyr Gln Glu Ser
 625 630 635 640
 Lys Glu Ala Asn Ser Ser Pro Gly Leu Gly Thr Pro Leu Lys Ser Pro
 645 650 655
 Ser Leu Ala Lys Lys Asp Ser Leu Ile Ser Glu Ser Glu Leu Ser Gln
 660 665 670
 Tyr Phe Ala His Asp Val Pro Gly Pro Leu Ser Asp Phe Met Val Ala
 675 680 685
 Gly Ser Glu Asp Glu Asp His Pro Gly Ser Gly Cys Ser Thr Ser Glu
 690 695 700
 Glu Gly Ser Leu Pro Pro Ser Thr Ser Ser Glu Pro Gly Lys Pro Arg
 705 710 715 720
 Ala Asn Ser Leu Val Thr Leu Gly Ser His Arg Ala Ser Gly Leu Phe
 725 730 735
 His Lys Gln Val Thr Val Ala Arg Gln Ala Ser Leu Pro Gly Ser Pro
 740 745 750
 Gln Ala Leu Arg Asn Pro Leu Leu Arg Gln Arg Lys Val Gly Cys Tyr
 755 760 765
 Asp Ala Asn Asp Ala Ser Asp Glu Glu Glu Phe Asp Arg Glu Gly Asp
 770 775 780
 Cys Ile Ser Leu Pro Gly Ala Leu Pro Gly Pro Ile Arg Pro Leu Ser
 785 790 795 800
 Glu Asp Asp Pro Arg Arg Val Ser Ile Ser Ser Ser Lys Gly Met Asp
 805 810 815
 Val His Asn Gln Glu Glu Arg Pro Arg Lys Thr Leu Val Ser Lys Ala
 820 825 830

Ile Ser Ala Pro Leu Leu Gly Ser Ser Val Asp Leu Glu Glu Ser Ile
 835 840 845

Pro Glu Gly Met Val Asp Ala Ala Ser Tyr Ala Ala Asn Leu Thr Asp
 850 855 860

Ser Ala Glu Ala Pro Lys Gly Ser Pro Gly Ser Trp Trp Lys Lys Glu
 865 870 875 880

Leu Ser Gly Ser Ser Ser Ala Pro Lys Leu Glu Tyr Thr Val Arg Thr
 885 890 895

Asp Thr Gln Ser Pro Thr Asn Thr Gly Ser Pro Ser Ser Pro Gln Gln
 900 905 910

Lys Ser Glu Gly Leu Gly Ser Arg His Arg Pro Val Ala Arg Val Ser
 915 920 925

Pro His Cys Lys Arg Ser Glu Ala Glu Ala Lys Pro Ser Gly Ser Gln
 930 935 940

Thr Val Asn Leu Thr Gly Arg Ala Asn Asp Pro Cys Asp Leu Asp Ser
 945 950 955 960

Arg Val Gln Ala Thr Ser Val Lys Val Thr Val Ala Gly Phe Gln Pro
 965 970 975

Gly Gly Ala Val Glu Lys Leu Cys Gln Glu Ser Leu Gly Lys Leu Thr
 980 985 990

Thr Gly Asp Ala Cys Val Ser Thr Ser Cys Glu Leu Ala Ser Ala Leu
 995 1000 1005

Ser His Leu Asp Ala Ser His Leu Thr Glu Asn Leu Pro Lys Ala Ala
 1010 1015 1020

Ser Glu Leu Gly Gln Gln Pro Met Thr Ser Ser Asp Leu Ile Ser Ser
 1025 1030 1035 1040

Pro Gly Lys Lys Gly Ala Ala His Pro Asp Pro Ser Lys Thr Ser Val
 1045 1050 1055

Asp Thr Gly Gln Val Ser Arg Pro Glu Asn Pro Ser Gln Pro Ala Ser
 1060 1065 1070

Pro Arg Val Thr Lys Cys Lys Ala Arg Ser Pro Val Arg Leu Pro His
 1075 1080 1085

Glu Gly Ser Pro Ser Pro Gly Glu Lys Ala Ala Ala Pro Pro Asp Tyr
 1090 1095 1100

Ser Lys Thr Arg Ser Ala Ser Glu Thr Ser Thr Pro His Asn Thr Arg
 1105 1110 1115 1120

Arg Val Ala Ala Leu Arg Gly Ala Gly Pro Gly Ala Glu Gly Met Thr
 1125 1130 1135

Pro Ala Gly Ala Val Leu Pro Gly Asp Pro Leu Thr Ser Gln Glu Gln
1140 1145 1150

Arg Gln Gly Ala Pro Gly Asn His Ser Lys Ala Leu Glu Met Thr Gly
1155 1160 1165

Ile His Ala Pro Glu Ser Ser Gln Glu Pro Ser Leu Leu Glu Gly Ala
1170 1175 1180

Asp Ser Val Ser Ser Arg Ala Pro Gln Ala Ser Leu Ser Met Leu Pro
1185 1190 1195 1200

Ser Thr Asp Asn Thr Lys Glu Ala Cys Gly His Val Ser Gly His Cys
1205 1210 1215

Cys Pro Gly Gly Ser Arg Glu Ser Pro Val Thr Asp Ile Asp Ser Phe
1220 1225 1230

Ile Lys Glu Leu Asp Ala Ser Ala Ala Arg Ser Pro Ser Ser Gln Thr
1235 1240 1245

Gly Asp Ser Gly Ser Gln Glu Gly Ser Ala Gln Gly His Pro Pro Ala
1250 1255 1260

Gly Ala Gly Gly Ser Ser Cys Arg Ala Glu Pro Val Pro Gly Gly
1265 1270 1275 1280

Gln Thr Ser Ser Pro Arg Arg Ala Trp Ala Ala Gly Ala Pro Ala Tyr
1285 1290 1295

Pro Gln Trp Ala Ser Gln Pro Ser Val Leu Asp Ser Ile Asn Pro Asp
1300 1305 1310

Lys His Phe Thr Val Asn Lys Asn Phe Leu Ser Asn Tyr Ser Arg Asn
1315 1320 1325

Phe Ser Ser Phe His Glu Asp Ser Thr Ser Leu Ser Gly Leu Gly Asp
1330 1335 1340

Ser Thr Glu Pro Ser Leu Ser Ser Met Tyr Gly Asp Ala Glu Asp Ser
1345 1350 1355 1360

Ser Ser Asp Pro Glu Ser Leu Thr Glu Ala Pro Arg Ala Ser Ala Arg
1365 1370 1375

Asp Gly Trp Ser Pro Pro Arg Ser Arg Val Ser Leu His Lys Glu Asp
1380 1385 1390

Pro Ser Glu Ser Glu Glu Gln Ile Glu Ile Cys Ser Thr Arg Gly
1395 1400 1405

Cys Pro Asn Pro Pro Ser Ser Pro Ala His Leu Pro Thr Gln Ala Ala
1410 1415 1420

Ile Cys Pro Ala Ser Ala Lys Val Leu Ser Leu Lys Tyr Ser Thr Pro
1425 1430 1435 1440

Arg Glu Ser Val Ala Ser Pro Arg Glu Lys Ala Ala Cys Leu Pro Gly
 1445 1450 1455

 Ser Tyr Thr Ser Gly Pro Asp Ser Ser Gln Pro Ser Ser Leu Leu Glu
 1460 1465 1470

 Met Ser Ser Gln Glu His Glu Thr His Ala Asp Ile Ser Thr Ser Gln
 1475 1480 1485

 Asn His Arg Pro Ser Cys Ala Glu Glu Thr Thr Glu Val Thr Ser Ala
 1490 1495 1500

 Ser Ser Ala Met Glu Asn Ser Pro Leu Ser Lys Val Ala Arg His Phe
 1505 1510 1515 1520

 His Ser Pro Pro Ile Ile Leu Ser Ser Pro Asn Met Val Asn Gly Leu
 1525 1530 1535

 Glu His Asp Leu Leu Asp Asp Glu Thr Leu Asn Gln Tyr Glu Thr Ser
 1540 1545 1550

 Ile Asn Ala Ala Ala Ser Leu Ser Ser Phe Ser Val Asp Val Pro Lys
 1555 1560 1565

 Asn Gly Glu Ser Val Leu Glu Asn Leu His Ile Ser Glu Ser Gln Asp
 1570 1575 1580

 Leu Asp Asp Leu Leu Gln Lys Pro Lys Met Ile Ala Arg Arg Pro Ile
 1585 1590 1595 1600

 Met Ala Trp Phe Lys Glu Ile Asn Lys His Asn Gln Gly Thr His Leu
 1605 1610 1615

 Arg Ser Lys Thr Glu Lys Glu Gln Pro Leu Met Pro Ala Arg Ser Pro
 1620 1625 1630

 Asp Ser Lys Ile Gln Met Val Ser Ser Ser Gln Lys Lys Gly Val Thr
 1635 1640 1645

 Val Pro His Ser Pro Pro Gln Pro Lys Thr Asn Leu Glu Asn Lys Asp
 1650 1655 1660

 Leu Ser Lys Lys Ser Pro Ala Glu Met Leu Leu Thr Asn Gly Gln Lys
 1665 1670 1675 1680

 Ala Lys Cys Gly Pro Lys Leu Lys Arg Leu Ser Leu Lys Gly Lys Ala
 1685 1690 1695

 Lys Val Asn Ser Glu Ala Pro Ala Ala Asn Ala Val Lys Ala Gly Gly
 1700 1705 1710

 Thr Asp His Arg Lys Pro Leu Ile Ser Pro Gln Thr Ser His Lys Thr
 1715 1720 1725

 Leu Ser Lys Ala Val Ser Gln Arg Leu His Val Ala Asp His Glu Asp
 1730 1735 1740

Pro Asp Arg Asn Thr Thr Ala Ala Pro Arg Ser Pro Gln Cys Val Leu
 1745 1750 1755 1760

Glu Ser Lys Pro Pro Leu Ala Thr Ser Gly Pro Leu Lys Pro Ser Val
 1765 1770 1775

Ser Asp Thr Ser Ile Arg Thr Phe Val Ser Pro Leu Thr Ser Pro Lys
 1780 1785 1790

Pro Val Pro Glu Gln Gly Met Trp Ser Arg Phe His Met Ala Val Leu
 1795 1800 1805

Ser Glu Pro Asp Arg Gly Cys Pro Thr Thr Pro Lys Ser Pro Lys Cys
 1810 1815 1820

Arg Ala Glu Gly Arg Ala Pro Arg Ala Asp Ser Gly Pro Val Ser Pro
 1825 1830 1835 1840

Ala Ala Ser Arg Asn Gly Met Ser Val Ala Gly Asn Arg Gln Ser Glu
 1845 1850 1855

Pro Arg Leu Ala Ser His Val Ala Ala Asp Thr Ala Gln Pro Arg Pro
 1860 1865 1870

Thr Gly Glu Lys Gly Gly Asn Ile Met Ala Ser Asp Arg Leu Glu Arg
 1875 1880 1885

Thr Asn Gln Leu Lys Ile Val Glu Ile Ser Ala Glu Ala Val Ser Glu
 1890 1895 1900

Thr Val Cys Gly Asn Lys Pro Ala Glu Ser Asp Arg Arg Gly Gly Cys
 1905 1910 1915 1920

Leu Ala Gln Gly Asn Cys Gln Glu Lys Ser Glu Ile Arg Leu Tyr Arg
 1925 1930 1935

Gln Val Ala Glu Ser Ser Thr Ser His Pro Ser Ser Leu Pro Ser His
 1940 1945 1950

Ala Ser Gln Ala Glu Gln Glu Met Ser Arg Ser Phe Ser Met Ala Lys
 1955 1960 1965

Leu Ala Ser Ser Ser Ser Leu Gln Thr Ala Ile Arg Lys Ala Glu
 1970 1975 1980

Tyr Ser Gln Gly Lys Ser Ser Leu Met Ser Asp Ser Arg Gly Val Pro
 1985 1990 1995 2000

Arg Asn Ser Ile Pro Gly Gly Pro Ser Gly Glu Asp His Leu Tyr Phe
 2005 2010 2015

Thr Pro Arg Pro Ala Thr Arg Thr Tyr Ser Met Pro Ala Gln Phe Ser
 2020 2025 2030

Ser His Phe Gly Arg Glu Gly His Pro Pro His Ser Leu Gly Arg Ser
 2035 2040 2045

Arg Asp Ser Gln Val Pro Val Thr Ser Ser Val Val Pro Glu Ala Lys
2050 2055 2060

Ala Ser Arg Gly Gly Leu Pro Ser Leu Ala Asn Gly Gln Gly Ile Tyr
2065 2070 2075 2080

Ser Val Lys Pro Leu Leu Asp Thr Ser Arg Asn Leu Pro Ala Thr Asp
2085 2090 2095

Glu Gly Asp Ile Ile Ser Val Gln Glu Thr Ser Cys Leu Val Thr Asp
2100 2105 2110

Lys Ile Lys Val Thr Arg Arg His Tyr Cys Tyr Glu Gln Asn Trp Pro
2115 2120 2125

His Glu Ser Thr Ser Phe Phe Ser Val Lys Gln Arg Ile Lys Ser Phe
2130 2135 2140

Glu Asn Leu Ala Asn Ala Asp Arg Pro Val Ala Lys Ser Gly Ala Ser
2145 2150 2155 2160

Pro Phe Leu Ser Val Ser Ser Lys Pro Pro Ile Gly Arg Arg Ser Ser
2165 2170 2175

Gly Ser Ile Val Ser Gly Ser Leu Gly His Pro Gly Asp Ala Ala Ala
2180 2185 2190

Arg Leu Leu Arg Arg Ser Leu Ser Ser Cys Ser Glu Asn Gln Ser Glu
2195 2200 2205

Ala Gly Thr Leu Leu Pro Gln Met Ala Lys Ser Pro Ser Ile Met Thr
2210 2215 2220

Leu Thr Ile Ser Arg Gln Asn Pro Pro Glu Thr Ser Ser Lys Gly Ser
2225 2230 2235 2240

Asp Ser Glu Leu Lys Lys Ser Leu Gly Pro Leu Gly Ile Pro Thr Pro
2245 2250 2255

Thr Met Thr Leu Ala Ser Pro Val Lys Arg Asn Lys Ser Ser Val Arg
2260 2265 2270

His Thr Gln Pro Ser Pro Val Ser Arg Ser Lys Leu Gln Glu Leu Arg
2275 2280 2285

Ala Leu Ser Met Pro Asp Leu Asp Lys Leu Cys Ser Glu Asp Tyr Ser
2290 2295 2300

Ala Gly Pro Ser Ala Val Leu Phe Lys Thr Glu Leu Glu Ile Thr Pro
2305 2310 2315 2320

Arg Arg Ser Pro Gly Pro Pro Ala Gly Gly Val Ser Cys Pro Glu Lys
2325 2330 2335

Gly Gly Asn Arg Ala Cys Pro Gly Gly Ser Gly Pro Lys Thr Ser Ala
2340 2345 2350

Ala Glu Thr Pro Ser Ser Ala Ser Asp Thr Gly Glu Ala Ala Gln Asp
2355 2360 2365

Leu Pro Phe Arg Arg Ser Trp Ser Val Lys Leu Asp Gln Leu Leu Val
2370 2375 2380

Ser Ala Gly Asp Gln Gln Arg Leu Gln Ser Val Leu Ser Ser Val Gly
2385 2390 2395 2400

Ser Lys Ser Thr Ile Leu Thr Leu Ile Gln Glu Ala Lys Ala Gln Ser
2405 2410 2415

Glu Asn Glu Glu Asp Val Cys Phe Ile Val Leu Asn Arg Lys Glu Gly
2420 2425 2430

Ser Gly Leu Gly Phe Ser Val Ala Gly Gly Thr Asp Val Glu Pro Lys
2435 2440 2445

Ser Ile Thr Val His Arg Val Phe Ser Gln Gly Ala Ala Ser Gln Glu
2450 2455 2460

Gly Thr Met Asn Arg Gly Asp Phe Leu Leu Ser Val Asn Gly Ala Ser
2465 2470 2475 2480

Leu Ala Gly Leu Ala His Gly Asn Val Leu Lys Val Leu His Gln Ala
2485 2490 2495

Gln Leu His Lys Asp Ala Leu Val Val Ile Lys Lys Gly Met Asp Gln
2500 2505 2510

Pro Arg Pro Ser Ala Arg Gln Glu Pro Pro Thr Ala Asn Gly Lys Gly
2515 2520 2525

Leu Leu Ser Arg Lys Thr Ile Pro Leu Glu Pro Gly Ile Gly Arg Ser
2530 2535 2540

Val Ala Val His Asp Ala Leu Cys Val Glu Val Leu Lys Thr Ser Ala
2545 2550 2555 2560

Gly Leu Gly Leu Ser Leu Asp Gly Gly Lys Ser Ser Val Thr Gly Asp
2565 2570 2575

Gly Pro Leu Val Ile Lys Arg Val Tyr Lys Gly Gly Ala Ala Glu Gln
2580 2585 2590

Ala Gly Ile Ile Glu Ala Gly Asp Glu Ile Leu Ala Ile Asn Gly Lys
2595 2600 2605

Pro Leu Val Gly Leu Met His Phe Asp Ala Trp Asn Ile Met Lys Ser
2610 2615 2620

Val Pro Glu Gly Pro Val Gln Leu Leu Ile Arg Lys His Arg Asn Ser
2625 2630 2635 2640

Ser

<210> 64
<211> 364
<212> PRT
<213> Mus musculus

<400> 64
Met Leu Arg Arg Phe Lys Gln Lys Ala His Leu Thr Tyr Asn Gly Asn
1 5 10 15

Ser Gly Asn Ser Ser Glu Pro Gly Glu Thr Pro Thr Leu Glu Leu Gly
20 25 30

Asp Gln Thr Ser Lys Lys Gly Lys Arg Thr Arg Lys Phe Gly Val Ile
35 40 45

Ser Arg Pro Ala Ile Ile Lys Ala Pro Glu Asp Ser Lys Ser Asn Ser
50 55 60

Gly Cys Asp Thr Ala Asp Asp Pro Ser Ser Glu Leu Glu Asn Gly Thr
65 70 75 80

Asp Ser Glu Leu Gly Asn Gly His Ala Phe Glu Leu Glu Asn Gly Pro
85 90 95

Asn Ser Leu Lys Asp Val Ala Gly Pro His Leu Glu Arg Ser Glu Ala
100 105 110

Asp Arg Glu Ala Glu Leu Arg Val Pro Lys Thr Glu Ala Pro Leu Ser
115 120 125

Asp Ser Asn Asp Lys Arg Arg Phe Ser Lys Thr Gly Lys Thr Asn Phe
130 135 140

Gln Ser Ser Asp Ser Leu Ala Arg Glu Glu Val Gly Arg Ile Trp Glu
145 150 155 160

Met Glu Leu Leu Lys Glu Ser Asp Gly Leu Gly Ile Gln Val Ser Gly
165 170 175

Gly Arg Gly Ser Lys Arg Ser Pro His Ala Ile Val Val Thr Gln Val
180 185 190

Lys Glu Gly Gly Ala Ala His Arg Asp Gly Arg Leu Ser Leu Gly Asp
195 200 205

Glu Leu Leu Val Ile Asn Gly His Leu Leu Val Gly Leu Ser His Glu
210 215 220

Glu Ala Val Ala Ile Leu Arg Ser Ala Thr Gly Met Val Gln Leu Val
225 230 235 240

Val Ala Ser Lys Met Leu Gly Ser Glu Ser Gln Asp Val Gly Ser
245 250 255

Ser Glu Glu Ser Lys Gly Asn Asn Leu Glu Ser Pro Lys Gln Gly Asn
260 265 270

Ser Lys Met Lys Leu Lys Ser Arg Leu Ser Gly Gly Val His Arg Leu
275 280 285

Glu Ser Val Glu Glu Tyr Asn Glu Leu Met Val Arg Asn Gly Asp Pro
290 295 300

Arg Ile Arg Met Leu Glu Val Ser Arg Asp Gly Arg Lys His Ser Leu
305 310 315 320

Pro Gln Leu Leu Asp Ser Thr Gly Thr Ser Gln Glu Tyr His Ile Val
325 330 335

Lys Lys Ser Thr Arg Ser Leu Ser Thr Thr His Val Glu Ser Pro Trp
340 345 350

Arg Leu Ile Arg Pro Ser Val Ile Ser Ile Ile Gly
355 360

<210> 65

<211> 51

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: LRRCT
Consensus Sequence

<400> 65

Asn Pro Phe Ile Cys Asp Cys Glu Leu Arg Trp Leu Leu Arg Trp Leu
1 5 10 15

Gln Ala Asn Arg His Leu Gln Asp Pro Val Asp Leu Arg Cys Ala Ser
20 25 30

Pro Glu Ser Leu Arg Gly Pro Leu Leu Leu Leu Pro Ser Ser Phe
35 40 45

Lys Cys Pro

50

<210> 66

<211> 51

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: LRRCT
Consensus Sequence

<400> 66

Asn Pro Phe Ile Cys Asp Cys Glu Leu Arg Trp Leu Leu Arg Trp Leu
1 5 10 15

Arg Glu Pro Arg Arg Leu Glu Asp Pro Glu Asp Leu Arg Cys Ala Ser

20

25

30

Pro Glu Ser Leu Arg Gly Pro Leu Leu Glu Leu Leu Pro Ser Asp Phe
35 40 45

Ser Cys Pro
50

<210> 67
<211> 124
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: CLECT
Consensus Sequence

<400> 67
Cys Pro Ser Gly Trp Val Ser Tyr Pro Gly Gly Lys Cys Tyr Lys Phe
1 5 10 15

Ser Thr Glu Lys Lys Thr Trp Ala Asp Ala Gln Ala Phe Cys Gln Ser
20 25 30

Leu Gly Ala His Leu Ala Ser Ile His Ser Glu Glu Asn Asp Phe
35 40 45

Leu Leu Ser Leu Leu Lys Asn Ser Asn Ser Asp Tyr Tyr Trp Ile Gly
50 55 60

Leu Ser Arg Pro Asp Ser Asn Gly Ser Trp Gln Trp Ser Asp Gly Ser
65 70 75 80

Gly Pro Val Asp Tyr Ser Asn Trp Ala Pro Gly Glu Pro Gly Gly Ser
85 90 95

Gly Asn Cys Val Val Leu Ser Thr Ser Gly Gly Lys Trp Asn Asp
100 105 110

Val Ser Cys Thr Ser Lys Leu Pro Phe Ile Cys Glu
115 120

<210> 68
<211> 114
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: CUB Consensus
Sequence

<400> 68
Cys Gly Gly Thr Leu Thr Ala Ser Ser Gly Thr Ile Thr Ser Pro Asn
1 5 10 15

Tyr Pro Asn Ser Tyr Pro Asn Asn Leu Asn Cys Val Trp Thr Ile Ser
20 25 30

Ala Pro Pro Gly Tyr Arg Ile Glu Leu Lys Phe Thr Asp Phe Asp Leu
35 40 45

Glu Ser Ser Asp Asn Cys Thr Tyr Asp Tyr Val Glu Ile Tyr Asp Gly
50 55 60

Pro Ser Thr Ser Ser Pro Leu Leu Gly Arg Phe Cys Gly Ser Glu Leu
65 70 75 80

Pro Pro Pro Ile Ile Ser Ser Ser Asn Ser Met Thr Val Thr Phe
85 90 95

Val Ser Asp Ser Ser Val Gln Lys Arg Gly Phe Ser Ala Arg Tyr Ser
100 105 110

Ala Val

<210> 69
<211> 110
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: CUB Consensus
Sequence

<400> 69
Cys Gly Gly Val Leu Thr Glu Ser Ser Gly Ser Ile Ser Ser Pro Asn
1 5 10 15

Tyr Pro Asn Asp Tyr Pro Pro Asn Lys Glu Cys Val Trp Thr Ile Arg
20 25 30

Ala Pro Pro Gly Tyr Arg Val Glu Leu Thr Phe Gln Asp Phe Asp Leu
35 40 45

Glu Asp His Thr Gly Cys Arg Tyr Asp Tyr Val Glu Ile Arg Asp Gly
50 55 60

Asp Gly Ser Ser Ser Pro Leu Leu Gly Lys Phe Cys Gly Ser Gly Pro
65 70 75 80

Pro Glu Asp Ile Val Ser Ser Ser Asn Arg Met Thr Ile Lys Phe Val
85 90 95

Ser Asp Ala Ser Val Ser Lys Arg Gly Phe Lys Ala Thr Tyr
100 105 110

<210> 70
<211> 107
<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Lectin C-type
domain Consensus Sequence

<400> 70

Glu Ser Lys Thr Trp Ala Glu Ala Gln Ala Ala Cys Gln Lys Leu Gly
1 5 10 15

Gly Gly Leu Val Ser Ile Gln Ser Ala Glu Glu Gln Asp Phe Leu Thr
20 25 30

Ser Leu Thr Lys Ala Ser Asn Ser Tyr Ala Trp Ile Gly Leu Thr Asp
35 40 45

Ile Asn Thr Glu Gly Thr Trp Val Trp Thr Asp Gly Ser Pro Val Asn
50 55 60

Tyr Thr Asn Trp Ala Pro Gly Glu Pro Asn Asn Arg Gly Asn Lys Glu
65 70 75 80

Asp Cys Val Glu Ile Tyr Thr Asp Gly Asn Lys Trp Asn Asp Glu Pro
85 90 95

Cys Gly Ser Lys Leu Pro Tyr Val Cys Glu Phe
100 105

<210> 71

<211> 304

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CNH domain
Consensus Sequence

<400> 71

Tyr Thr Ala Lys Cys Asn His Pro Ile Thr Cys Asp Ala Lys Asn Leu
1 5 10 15

Leu Val Gly Thr Glu Glu Gly Leu Tyr Val Leu Asn Arg Ser Asp Gln
20 25 30

Gly Gly Thr Leu Glu Lys Ile Ile Ser Arg Arg Ser Val Thr Gln Ile
35 40 45

Trp Val Leu Glu Glu Asn Asn Val Leu Leu Met Ile Ser Gly Lys Lys
50 55 60 80

Pro Tyr Leu Tyr Ala His Pro Leu Ser Gly Leu Arg Glu Lys Asp Ala
65 70 75 80

Leu Gly Ser Ala Arg Leu Val Ile Arg Lys Asn Val Trp Val Lys Ile
85 90 95

Glu	Asp	Val	Lys	Gly	Cys	His	Leu	Phe	Ala	Val	Val	Asn	Gly	Lys	Arg
100							105					110			
Ile	Leu	Phe	Leu	Cys	Ala	Ala	Leu	Pro	Ser	Ser	Val	Gln	Leu	Leu	Gln
115							120					125			
Trp	Tyr	Asn	Pro	Leu	Lys	Lys	Phe	Lys	Leu	Phe	Lys	Ser	Lys	Phe	Leu
130					135				140						
Lys	Lys	Leu	Ile	Val	Pro	Val	Pro	Leu	Phe	Val	Leu	Leu	Thr	Ser	Ser
145					150				155				160		
Ser	Phe	Glu	Leu	Pro	Lys	Ile	Cys	Ile	Gly	Val	Asp	Lys	Asn	Gly	Phe
	165						170					175			
Asp	Val	Val	Gln	Phe	His	Gln	Thr	Ser	Leu	Val	Ser	Lys	Glu	Asp	Leu
	180						185					190			
Ser	Leu	Pro	Asn	Leu	Asn	Glu	Glu	Thr	Ser	Lys	Lys	Pro	Ile	Pro	Val
	195					200				205					
Ile	Gln	Val	Pro	Gln	Ser	Asp	Asp	Glu	Leu	Leu	Leu	Cys	Phe	Asp	Glu
	210					215				220					
Phe	Gly	Val	Phe	Val	Asn	Leu	Gln	Gly	Met	Arg	Arg	Ser	Arg	Lys	Pro
	225					230			235			240			
Ile	Leu	Thr	Trp	Glu	Phe	Met	Pro	Glu	Tyr	Phe	Ala	Tyr	His	Glu	Pro
						245		250				255			
Tyr	Leu	Leu	Ala	Phe	His	Ser	Asn	Gly	Ile	Glu	Ile	Arg	Ser	Ile	Glu
						260		265			270				
Thr	Gly	Glu	Leu	Leu	Gln	Glu	Leu	Ala	Asp	Arg	Glu	Ala	Arg	Lys	Ile
						275		280			285				
Arg	Val	Leu	Gly	Ser	Ser	Asp	Arg	Lys	Ile	Leu	Val	Ser	Ser	Ser	Pro
						290		295			300				

<210> 72
<211> 256
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Serine/Threonine protein kinase Consensus
Sequence

<400> 72
Tyr Glu Leu Leu Glu Val Leu Gly Lys Gly Ala Phe Gly Lys Val Tyr
1 5 10 15

Leu	Ala	Arg	Asp	Lys	Lys	Thr	Gly	Lys	Leu	Val	Ala	Ile	Lys	Val	Ile
				20				25					30		
Lys	Lys	Glu	Lys	Leu	Lys	Lys	Lys	Arg	Glu	Arg	Ile	Leu	Arg	Glu	
		35						40					45		
Ile	Lys	Ile	Leu	Lys	Lys	Leu	Asp	His	Pro	Asn	Ile	Val	Lys	Leu	Tyr
		50					55					60			
Asp	Val	Phe	Glu	Asp	Asp	Asp	Lys	Leu	Tyr	Leu	Val	Met	Glu	Tyr	Cys
	65						70			75			80		
Glu	Gly	Gly	Asp	Leu	Phe	Asp	Leu	Leu	Lys	Lys	Arg	Gly	Arg	Leu	Ser
			85						90				95		
Glu	Asp	Glu	Ala	Arg	Phe	Tyr	Ala	Arg	Gln	Ile	Leu	Ser	Ala	Leu	Glu
		100						105					110		
Tyr	Leu	His	Ser	Gln	Gly	Ile	Ile	His	Arg	Asp	Leu	Lys	Pro	Glu	Asn
		115						120					125		
Ile	Leu	Leu	Asp	Ser	Asp	Gly	His	Val	Lys	Leu	Ala	Asp	Phe	Gly	Leu
		130					135					140			
Ala	Lys	Gln	Leu	Asp	Ser	Gly	Gly	Thr	Leu	Leu	Thr	Thr	Phe	Val	Gly
	145					150			155				160		
Thr	Pro	Glu	Tyr	Met	Ala	Pro	Glu	Val	Leu	Leu	Gly	Lys	Gly	Tyr	Gly
		165					170					175			
Lys	Ala	Val	Asp	Ile	Trp	Ser	Leu	Gly	Val	Ile	Leu	Tyr	Glu	Leu	Leu
		180						185				190			
Thr	Gly	Lys	Pro	Pro	Phe	Pro	Gly	Asp	Asp	Gln	Leu	Leu	Ala	Leu	Phe
		195					200					205			
Lys	Lys	Ile	Gly	Lys	Pro	Pro	Pro	Phe	Pro	Pro	Pro	Glu	Trp	Lys	
		210					215					220			
Ile	Ser	Pro	Glu	Ala	Lys	Asp	Leu	Ile	Lys	Lys	Leu	Leu	Val	Lys	Asp
		225					230				235			240	
Pro	Glu	Lys	Arg	Leu	Thr	Ala	Glu	Ala	Leu	Glu	His	Pro	Phe	Phe	
			245					250				255			

<210> 73

<211> 301

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: CNH domain
Consensus Sequence

<400> 73
 Asn Thr Ala Lys Trp Asn His Pro Ile Thr Cys Asp Ala Lys Ile Leu
 1 5 10 15
 Leu Val Gly Thr Glu Glu Gly Leu Tyr Val Leu Asn Ile Ser Asp Gln
 20 25 30
 His Gly Thr Leu Glu Lys Leu Ile Gly Arg Arg Ser Val Thr Gln Ile
 35 40 45
 Trp Val Leu Glu Glu Asn Asn Val Leu Leu Met Ile Ser Gly Lys Lys
 50 55 60
 Pro Gln Leu Tyr Ser His Pro Leu Ser Ala Leu Thr Glu Lys Asp Ala
 65 70 75 80
 Leu Gly Ser Ala Arg Leu Val Ile Arg Lys Asn Val Leu Thr Lys Ile
 85 90 95
 Pro Asp Val Lys Gly Cys His Leu Cys Ala Val Val Asn Gly Lys Arg
 100 105 110
 Ile Leu Phe Leu Cys His Ala Leu Gln Ser Ser Val Val Leu Leu Gln
 115 120 125
 Trp Tyr Asn Pro Leu Lys Lys Phe Lys Leu Phe Lys Ser Lys Phe Leu
 130 135 140
 Phe Pro Leu Ile Ser Pro Val Pro Val Phe Val Glu Leu Val Ser Ser
 145 150 155 160
 Ser Phe Glu Leu Pro Gly Ile Cys Ile Gly Ser Asp Lys Asn Gly Gly
 165 170 175
 Asp Val Val Gln Phe His Gln Ser Leu Val Ser Lys Glu Asp Leu Ser
 180 185 190
 Leu Pro Phe Leu Ser Glu Glu Thr Ser Ser Lys Pro Ile Ser Val Val
 195 200 205
 Gln Val Pro Ala Asp Glu Leu Leu Cys Tyr Asp Glu Phe Gly Val
 210 215 220
 Phe Val Asn Leu Tyr Gly Met Arg Arg Ser Arg Asn Pro Ile Leu His
 225 230 235 240
 Trp Glu Phe Met Pro Glu Ser Phe Ala Tyr His Ser Pro Tyr Leu Leu
 245 250 255
 Ala Phe His Asp Asn Gly Ile Glu Ile Arg Ser Ile Lys Thr Gly Glu
 260 265 270
 Leu Leu Gln Glu Leu Ala Asp Arg Lys Thr Arg Lys Ile Arg Leu Leu
 275 280 285
 Gly Ser Ser Asp Arg Lys Ile Leu Leu Ser Ser Ser Pro

290

295

300

<210> 74
<211> 256
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Protein kinase
domain Consensus Sequence

<400> 74
Tyr Glu Leu Gly Glu Lys Leu Gly Ser Gly Ala Phe Gly Lys Val Tyr
1 5 10 15

Lys Gly Lys His Lys Asp Thr Gly Glu Ile Val Ala Ile Lys Ile Leu
20 25 30

Lys Lys Arg Ser Leu Ser Glu Lys Lys Arg Phe Leu Arg Glu Ile
35 40 45

Gln Ile Leu Arg Arg Leu Ser His Pro Asn Ile Val Arg Leu Leu Gly
50 55 60

Val Phe Glu Glu Asp Asp His Leu Tyr Leu Val Met Glu Tyr Met Glu
65 70 75 80

Gly Gly Asp Leu Phe Asp Tyr Leu Arg Arg Asn Gly Leu Leu Ser
85 90 95

Glu Lys Glu Ala Lys Lys Ile Ala Leu Gln Ile Leu Arg Gly Leu Glu
100 105 110

Tyr Leu His Ser Arg Gly Ile Val His Arg Asp Leu Lys Pro Glu Asn
115 120 125

Ile Leu Leu Asp Glu Asn Gly Thr Val Lys Ile Ala Asp Phe Gly Leu
130 135 140

Ala Arg Lys Leu Glu Ser Ser Tyr Glu Lys Leu Thr Thr Phe Val
145 150 155 160

Gly Thr Pro Glu Tyr Met Ala Pro Glu Val Leu Glu Gly Arg Gly Tyr
165 170 175

Ser Ser Lys Val Asp Val Trp Ser Leu Gly Val Ile Leu Tyr Glu Leu
180 185 190

Leu Thr Gly Lys Leu Pro Phe Pro Gly Ile Asp Pro Leu Glu Glu Leu
195 200 205

Phe Arg Ile Lys Glu Arg Pro Arg Leu Arg Leu Pro Leu Pro Pro Asn
210 215 220

Cys Ser Glu Glu Leu Lys Asp Leu Ile Lys Lys Cys Leu Asn Lys Asp
225 230 235 240

Pro Glu Lys Arg Pro Thr Ala Lys Glu Ile Leu Asn His Pro Trp Phe
245 250 255

<210> 75
<211> 258
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Tyrosine
Kinase Consensus Sequence

<400> 75
Leu Thr Leu Gly Lys Lys Leu Gly Glu Gly Ala Phe Gly Glu Val Tyr
1 5 10 15

Lys Gly Thr Leu Lys Gly Lys Gly Val Glu Val Glu Val Ala Val
20 25 30

Lys Thr Leu Lys Glu Asp Ala Ser Glu Gln Gln Ile Glu Glu Phe Leu
35 40 45

Arg Glu Ala Arg Leu Met Arg Lys Leu Asp His Pro Asn Ile Val Lys
50 55 60

Leu Leu Gly Val Cys Thr Glu Glu Pro Leu Met Ile Val Met Glu
65 70 75 80

Tyr Met Glu Gly Gly Asp Leu Leu Asp Tyr Leu Arg Lys Asn Arg Pro
85 90 95

Lys Glu Leu Ser Leu Ser Asp Leu Leu Ser Phe Ala Leu Gln Ile Ala
100 105 110

Arg Gly Met Glu Tyr Leu Glu Ser Lys Asn Phe Val His Arg Asp Leu
115 120 125

Ala Ala Arg Asn Cys Leu Val Gly Glu Asn Lys Thr Val Lys Ile Ala
130 135 140

Asp Phe Gly Leu Ala Arg Asp Leu Tyr Asp Asp Asp Tyr Tyr Arg Lys
145 150 155 160

Lys Lys Ser Pro Arg Leu Pro Ile Arg Trp Met Ala Pro Glu Ser Leu
165 170 175

Lys Asp Gly Lys Phe Thr Ser Lys Ser Asp Val Trp Ser Phe Gly Val
180 185 190

Leu Leu Trp Glu Ile Phe Thr Leu Gly Glu Ser Pro Tyr Pro Gly Met
195 200 205

Ser	Asn	Glu	Glu	Val	Leu	Glu	Tyr	Leu	Lys	Lys	Gly	Tyr	Arg	Leu	Pro
210						215					220				
Gln	Pro	Pro	Asn	Cys	Pro	Asp	Glu	Ile	Tyr	Asp	Leu	Met	Leu	Gln	Cys
225						230				235					240
Trp	Ala	Glu	Asp	Pro	Glu	Asp	Arg	Pro	Thr	Phe	Ser	Glu	Leu	Val	Glu
									245				250		255
Arg	Leu														

<210>	76														
<211>	430														
<212>	PRT														
<213>	Artificial Sequence														
<220>															
<223>	Description of Artificial Sequence:	Sema domain													
	Consensus Sequence														
<400>	76														
Leu	Gln	Asn	Leu	Leu	Leu	Asp	Glu	Asp	Asn	Gly	Thr	Leu	Tyr	Val	Gly
1									10					15	
Ala	Arg	Asn	Arg	Leu	Tyr	Val	Leu	Ser	Leu	Asn	Leu	Ile	Ser	Glu	Ala
				20				25					30		
Glu	Val	Lys	Thr	Gly	Pro	Val	Leu	Ser	Ser	Pro	Asp	Cys	Glu	Glu	Cys
								35				40		45	
Val	Ser	Lys	Gly	Lys	Asp	Pro	Pro	Thr	Asp	Cys	Val	Asn	Phe	Ile	Arg
								50			55		60		
Leu	Leu	Leu	Asp	Tyr	Asn	Ala	Asp	His	Leu	Leu	Val	Cys	Gly	Thr	Asn
								65			70		75		80
Ala	Phe	Gln	Pro	Val	Cys	Arg	Leu	Ile	Asn	Leu	Gly	Asn	Leu	Asp	Arg
								85			90			95	
Leu	Glu	Val	Gly	Arg	Glu	Ser	Gly	Arg	Gly	Arg	Cys	Pro	Phe	Asp	Pro
								100			105			110	
Gln	His	Asn	Ser	Thr	Ala	Val	Leu	Val	Asp	Gly	Glu	Leu	Tyr	Val	Gly
								115			120			125	
Thr	Val	Ala	Asp	Phe	Ser	Gly	Ser	Asp	Pro	Ala	Ile	Tyr	Arg	Ser	Leu
								130			135		140		
Ser	Val	Arg	Arg	Leu	Lys	Gly	Thr	Ser	Gly	Pro	Ser	Leu	Arg	Thr	Val
								145			150		155		160
Leu	Tyr	Asp	Ser	Arg	Trp	Leu	Asn	Glu	Pro	Asn	Phe	Val	Tyr	Ala	Phe
								165			170			175	
Glu	Ser	Gly	Asp	Phe	Val	Tyr	Phe	Phe	Phe	Arg	Glu	Thr	Ala	Val	Glu

180

185

190

Asp Glu Asn Cys Gly Lys Ala Val Val Ser Arg Val Ala Arg Val Cys
 195 200 205

Lys Asn Asp Val Gly Gly Pro Arg Ser Leu Ser Lys Lys Trp Thr Ser
 210 215 220

Phe Leu Lys Ala Arg Leu Glu Cys Ser Val Pro Gly Glu Phe Pro Phe
 225 230 235 240

Tyr Phe Asn Glu Leu Gln Ala Ala Phe Leu Leu Pro Ala Gly Ser Glu
 245 250 255

Ser Asp Asp Val Leu Tyr Gly Val Phe Ser Thr Ser Ser Asn Pro Ile
 260 265 270

Pro Gly Ser Ala Val Cys Ala Phe Ser Leu Ser Asp Ile Asn Ala Val
 275 280 285

Phe Asn Glu Pro Phe Lys Glu Cys Glu Thr Gly Asn Ser Gln Trp Leu
 290 295 300

Pro Tyr Pro Arg Gly Leu Val Pro Phe Pro Arg Pro Gly Thr Cys Pro
 305 310 315 320

Asn Thr Pro Leu Ser Ser Lys Asp Leu Pro Asp Asp Val Leu Asn Phe
 325 330 335

Ile Lys Thr His Pro Leu Met Asp Glu Val Val Gln Pro Leu Thr Gly
 340 345 350

Arg Pro Leu Phe Val Lys Thr Asp Ser Asn Tyr Leu Leu Thr Ser Ile
 355 360 365

Ala Val Asp Arg Val Arg Thr Asp Gly Gly Asn Tyr Thr Val Leu Phe
 370 375 380

Leu Gly Thr Ser Asp Gly Arg Ile Leu Lys Val Val Leu Ser Arg Ser
 385 390 395 400

Ser Ser Ser Ser Glu Ser Val Val Leu Glu Glu Ile Ser Val Phe Asp
 405 410 415

Pro Gly Ser Pro Val Ser Asp Leu Val Leu Ser Pro Lys Lys
 420 425 430

<210> 77

<211> 433

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Sema domain
Consensus Sequence

<400> 77

Phe Val Thr Leu Leu Leu Asp Glu Asp Arg Gly Arg Leu Tyr Val Gly
1 5 10 15

Ala Arg Asn Arg Val Tyr Val Leu Asn Leu Glu Asp Leu Ser Glu Val
20 25 30

Leu Asn Leu Lys Thr Gly Trp Pro Gly Ser Cys Glu Thr Cys Glu Glu
35 40 45

Cys Asn Met Lys Gly Lys Ser Pro Leu Thr Glu Cys Thr Asn Phe Ile
50 55 60

Arg Val Leu Gln Ala Tyr Asn Asp Thr His Leu Tyr Val Cys Gly Thr
65 70 75 80

Asn Ala Phe Gln Pro Val Cys Thr Leu Ile Asn Leu Gly Asp Leu Phe
85 90 95

Ser Leu Asp Val Asp Asn Glu Glu Asp Gly Cys Gly Asp Cys Pro Tyr
100 105 110

Asp Pro Leu Gly Asn Thr Thr Ser Val Leu Val Gln Gly Gly Glu Leu
115 120 125

Tyr Ser Gly Thr Val Ile Asp Phe Ser Gly Arg Asp Pro Ser Ile Arg
130 135 140

Arg Leu Leu Gly Ser His Asp Gly Leu Arg Thr Glu Phe His Asp Ser
145 150 155 160

Lys Trp Leu Asn Leu Pro Asn Phe Val Asp Ser Tyr Pro Ile His Tyr
165 170 175

Val His Ser Phe Ser Asp Asp Lys Val Tyr Phe Phe Arg Glu Thr
180 185 190

Ala Val Glu Asp Ser Asn Cys Lys Thr Ile His Ser Arg Val Ala Arg
195 200 205

Val Cys Lys Asn Asp Pro Gly Gly Arg Ser Tyr Leu Glu Leu Asn Lys
210 215 220

Trp Thr Thr Phe Leu Lys Ala Arg Leu Asn Cys Ser Ile Pro Gly Glu
225 230 235 240

Gly Thr Pro Phe Tyr Phe Asn Glu Leu Gln Ala Ala Phe Val Leu Pro
245 250 255

Thr Gly Ala Asp Thr Asp Pro Val Leu Tyr Gly Val Phe Thr Thr Ser
260 265 270

Ser Asn Ser Ser Ala Gly Ser Ala Val Cys Ala Phe Ser Met Ser Asp
275 280 285

Ile Asn Gln Val Phe Glu Gly Pro Phe Lys His Gln Ser Pro Asn Ser
290 295 300

Lys	Trp	Leu	Pro	Tyr	Arg	Gly	Lys	Val	Pro	Gln	Pro	Arg	Pro	Gly	Gln
305															320
Cys	Pro	Asn	Ala	Ser	Gly	Leu	Asn	Leu	Pro	Asp	Asp	Thr	Leu	Asn	Phe
	325														335
Ile	Arg	Cys	His	Pro	Leu	Met	Asp	Glu	Val	Val	Pro	Pro	Leu	His	Asn
		340													350
Val	Pro	Leu	Phe	Val	Gly	Gln	Ser	Gly	Asn	Tyr	Arg	Leu	Thr	Ser	Ile
		355													365
Ala	Val	Asp	Arg	Val	Arg	Ala	Gly	Asp	Gly	Gln	Ile	Tyr	Thr	Val	Leu
		370													380
Phe	Leu	Gly	Thr	Asp	Asp	Gly	Arg	Val	Leu	Lys	Gln	Val	Val	Leu	Ser
		385													400
Arg	Ser	Ser	Ser	Ala	Ser	Tyr	Leu	Val	Val	Val	Leu	Glu	Glu	Ser	Leu
															415
Val	Phe	Pro	Asp	Gly	Glu	Pro	Val	Gln	Arg	Met	Val	Ile	Ser	Ser	Lys
		420													430

Asn

<210> 78
<211> 85
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: TIG domain
Consensus Sequence

<400> 78
Pro Val Ile Thr Ser Ile Ser Pro Ser Ser Gly Pro Leu Ser Gly Gly
1 5 10 15

Thr Glu Ile Thr Ile Thr Gly Ser Asn Leu Gly Ser Gly Glu Asp Ile
20 25 30

Lys Val Thr Phe Gly Gly Thr Glu Cys Asp Val Val Ser Gln Glu Ala
35 40 45

Ser Gln Ile Val Cys Lys Thr Pro Pro Tyr Ala Asn Gly Gly Pro Gln
50 55 60

Pro Val Thr Val Ser Leu Asp Gly Gly Leu Ser Ser Ser Pro Val
65 70 75 80

Thr Phe Thr Tyr Val
85

<210> 79
<211> 85
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: TIG Consensus Sequence

<400> 79
Pro Val Ile Thr Ser Ile Ser Pro Ser Ser Gly Pro Leu Ser Gly Gly
1 5 10 15

Thr Glu Ile Thr Ile Thr Gly Ser Asn Leu Gly Ser Gly Glu Asp Ile
20 25 30

Lys Val Thr Phe Gly Gly Thr Glu Cys Asp Val Val Ser Gln Glu Ala
35 40 45

Ser Gln Ile Val Cys Lys Thr Pro Pro Tyr Ala Asn Gly Gly Pro Gln
50 55 60

Pro Val Thr Val Ser Leu Asp Gly Gly Leu Ser Ser Ser Pro Val
65 70 75 80

Thr Phe Thr Tyr Val
85

<210> 80
<211> 85
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: TIG Consensus Sequence

<400> 80
Pro Val Ile Thr Ser Ile Ser Pro Ser Ser Gly Pro Leu Ser Gly Gly
1 5 10 15

Thr Glu Ile Thr Ile Thr Gly Ser Asn Leu Gly Ser Gly Glu Asp Ile
20 25 30

Lys Val Thr Phe Gly Gly Thr Glu Cys Asp Val Val Ser Gln Glu Ala
35 40 45

Ser Gln Ile Val Cys Lys Thr Pro Pro Tyr Ala Asn Gly Gly Pro Gln
50 55 60

Pro Val Thr Val Ser Leu Asp Gly Gly Leu Ser Ser Ser Pro Val
65 70 75 80

Thr Phe Thr Tyr Val
85

<210> 81
<211> 47
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: PSI domain
Consensus Sequence

<400> 81
Arg Cys Ser Ala Tyr Thr Ser Cys Ser Glu Cys Leu Leu Ala Arg Asp
1 5 10 15

Pro Tyr Cys Ala Trp Cys Ser Ser Gln Gly Arg Cys Thr Ser Gly Glu
20 25 30

Arg Cys Asp Ser Leu Arg Gln Asn Trp Ser Ser Gly Gln Cys Pro
35 40 45

<210> 82
<211> 93
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: IPT Consensus
Sequence

<400> 82
Asp Pro Val Ile Thr Arg Ile Ser Pro Asn Ser Gly Pro Leu Ser Gly
1 5 10 15

Gly Thr Arg Ile Thr Leu Cys Gly Lys Asn Leu Asp Ser Ile Ser Val
20 25 30

Val Phe Val Glu Val Gly Val Gly Glu Val Pro Cys Thr Phe Leu Pro
35 40 45

Ser Asp Val Ser Gln Thr Ala Ile Val Cys Lys Thr Pro Pro Tyr His
50 55 60

Asn Ile Pro Gly Ser Val Pro Val Arg Val Glu Val Gly Leu Arg Asn
65 70 75 80

Gly Gly Val Pro Gly Glu Pro Ser Pro Phe Thr Tyr Val
85 90

<210> 83
<211> 254
<212> PRT
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: 7
transmembrane receptor Consensus Sequence

<400> 83

Gly Asn Leu Leu Val Ile Leu Val Ile Leu Arg Thr Lys Lys Leu Arg
1 5 10 15

Thr Pro Thr Asn Ile Phe Leu Leu Asn Leu Ala Val Ala Asp Leu Leu
20 25 30

Phe Leu Leu Thr Leu Pro Pro Trp Ala Leu Tyr Tyr Leu Val Gly Gly
35 40 45

Asp Trp Val Phe Gly Asp Ala Leu Cys Lys Leu Val Gly Ala Leu Phe
50 55 60

Val Val Asn Gly Tyr Ala Ser Ile Leu Leu Leu Thr Ala Ile Ser Ile
65 70 75 80

Asp Arg Tyr Leu Ala Ile Val His Pro Leu Arg Tyr Arg Arg Ile Arg
85 90 95

Thr Pro Arg Arg Ala Lys Val Leu Ile Leu Leu Val Trp Val Leu Ala
100 105 110

Leu Leu Leu Ser Leu Pro Pro Leu Leu Phe Ser Trp Leu Arg Thr Val
115 120 125

Glu Glu Gly Asn Thr Thr Val Cys Leu Ile Asp Phe Pro Glu Glu Ser
130 135 140

Val Lys Arg Ser Tyr Val Leu Leu Ser Thr Leu Val Gly Phe Val Leu
145 150 155 160

Pro Leu Leu Val Ile Leu Val Cys Tyr Thr Arg Ile Leu Arg Thr Leu
165 170 175

Arg Lys Arg Ala Arg Ser Gln Arg Ser Leu Lys Arg Arg Ser Ser Ser
180 185 190

Glu Arg Lys Ala Ala Lys Met Leu Leu Val Val Val Val Phe Val
195 200 205

Leu Cys Trp Leu Pro Tyr His Ile Val Leu Leu Leu Asp Ser Leu Cys
210 215 220

Leu Leu Ser Ile Trp Arg Val Leu Pro Thr Ala Leu Leu Ile Thr Leu
225 230 235 240

Trp Leu Ala Tyr Val Asn Ser Cys Leu Asn Pro Ile Ile Tyr
245 250

<210> 84

<211> 402

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: ANF receptor
Consensus Sequence

<400> 84

Gly	Leu	Ile	Asn	Tyr	Ala	Val	Arg	Gly	Ile	Thr	Arg	Leu	Glu	Ala	Met
1									10						15
Leu Gly Ala Phe Asp Arg Ile Asn Ala Asp Pro Ala Leu Leu Pro Gly															
						20			25						30
Leu Ala Leu Gly Leu Ala Ile Leu Asp Ile Asn Ser Leu Arg Asn Val															
						35			40						45
Ala Leu Glu Gln Ser Phe Thr Phe Val Tyr Gly Leu Leu Ile Lys Cys															
						50			55						60
Asp Cys Ser Ser Val Arg Cys Ala Gly Gly Asp Leu Ala Leu Thr His															
						65			70						80
Gly Val Ala Gly Val Ile Gly Pro Ser Cys Ser Ser Ser Ala Ile Gln															
						85			90						95
Val Ala Asn Leu Ala Ser Leu Leu Asn Ile Pro Met Ile Ser Tyr Gly															
						100			105						110
Ser Thr Ala Pro Glu Leu Ser Asp Lys Thr Arg Tyr Pro Thr Phe Ser															
						115			120						125
Arg Thr Ile Pro Ser Asp Ala Phe Gln Gly Leu Ala Met Val Asp Ile															
						130			135						140
Phe Lys His Phe Asn Trp Asn Tyr Val Ser Val Val Tyr Ser Asp Gly															
						145			150						160
Thr Tyr Gly Glu Glu Gly Cys Glu Ala Phe Ile Glu Ala Leu Arg Ala															
						165			170						175
Arg Gly Gly Cys Ile Ala Leu Ser Val Lys Ile Gly Glu Phe Asp Arg															
						180			185						190
Gly Asp Glu Glu Asp Phe Asp Lys Leu Leu Arg Glu Leu Lys Arg Arg															
						195			200						205
Ala Arg Val Val Val Met Cys Gly His Gly Glu Thr Leu Arg Glu Leu															
						210			215						220
Leu Glu Ala Ala Leu Arg Leu Gly Leu Thr Gly Glu Asp Tyr Val Phe															
						225			230						240
Ile Ser Asp Asp Leu Phe Asn Lys Ser Leu Pro Ala Glu Pro Gly Ala															
						245			250						255
Pro Gly Ala Ile Glu Leu Ala Asn Ala Ser Met Leu Arg Phe Ala Tyr															
						260			265						270

Tyr Phe Val Leu Val Leu Thr Leu Asn Asn Pro Arg Asn Pro Trp Phe
 275 280 285

Leu Glu Phe Trp Lys Glu Asn Phe Ile Cys Ala Leu Gln Asp Phe Leu
 290 295 300

Thr Leu Glu Pro Tyr Glu Gln Glu Gly Lys Ala Gly Phe Val Tyr Asp
 305 310 315 320

Ala Val Tyr Leu Tyr Ala His Ala Leu His Asn Thr Thr Leu Ala Leu
 325 330 335

Gly Gly Ser Trp Val Asp Gly Glu Lys Leu Val Gln His Leu Arg Asn
 340 345 350

Leu Thr Phe Glu Gly Val Thr Gly Pro Val Thr Phe Asp Glu Asn Gly
 355 360 365

Asp Arg Asp Gly Asp Tyr Val Leu Leu Asp Thr Gln Asn Thr Glu Thr
 370 375 380

Gly Gln Leu Lys Val Thr Gly Thr Tyr Asp Gly Val Gly Lys Trp Thr
 385 390 395 400

Glu Pro

<210> 85
<211> 256
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: 7
transmembrane receptor (metabotropic E family)
Consensus Sequence

<400> 85
Leu Gly Ile Val Leu Val Ala Leu Ala Val Leu Gly Ile Val Leu Thr
1 5 10 15

Leu Phe Val Leu Val Val Phe Val Lys His Arg Asp Thr Pro Ile Val
20 25 30

Lys Ala Ser Asn Arg Glu Leu Ser Tyr Leu Leu Leu Ile Gly Leu Ile
 35 40 45

Leu Cys Tyr Leu Cys Ser Phe Leu Phe Ile Gly Lys Pro Ser Glu Thr

Ser Cys Ile Leu Arg Arg Ile Leu Phe Gly Leu Gly Phe Thr Leu Cys

Tyr Ser Ala Leu Leu Ala Lys Thr Asn Arg Val Leu Arg Ile Phe Arg

Ala Lys Lys Pro Gly Ser Gly Lys Pro Lys Phe Ile Ser Pro Trp Ala
 100 105 110

 Gln Val Leu Ile Val Leu Ile Leu Val Leu Ile Gln Val Ile Ile Cys
 115 120 125

 Val Ile Trp Leu Val Val Glu Pro Pro Arg Pro Thr Ile Asp Ile Tyr
 130 135 140

 Ser Glu Lys Glu Lys Ile Ile Leu Glu Cys Asn Lys Gly Ser Met Val
 145 150 155 160

 Ala Phe Val Val Leu Gly Tyr Asp Gly Leu Leu Ala Val Leu Cys
 165 170 175

 Thr Phe Leu Ala Phe Leu Thr Arg Asn Leu Pro Glu Asn Phe Asn Glu
 180 185 190

 Ala Lys Phe Ile Gly Phe Ser Met Leu Thr Phe Cys Ile Val Trp Val
 195 200 205

 Ala Phe Ile Pro Ile Tyr Leu Ser Thr Pro Gly Lys Val Gln Val Ala
 210 215 220

 Val Glu Ile Phe Ser Ile Leu Ala Ser Ser Thr Val Leu Leu Gly Cys
 225 230 235 240

 Leu Phe Val Pro Lys Cys Tyr Ile Ile Leu Phe Arg Pro Glu Lys Asn
 245 250 255

<210> 86
 <211> 86
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: PDZ domain
 Consensus Sequence

<400> 86
 Glu Pro Arg Leu Val Glu Leu Glu Lys Gly Gly Gly Gly Leu Gly Phe
 1 5 10 15

 Ser Leu Val Gly Gly Lys Asp Ser Gly Asp Gly Gly Val Val Val Ser
 20 25 30

 Ser Val Val Pro Gly Ser Pro Ala Ala Lys Ala Gly Leu Lys Pro Gly
 35 40 45

 Asp Val Ile Leu Glu Val Asn Gly Thr Ser Val Glu Gly Leu Thr His
 50 55 60

 Leu Glu Ala Val Asp Leu Leu Lys Glu Ala Gly Gly Lys Val Thr Leu

65 70 75 80

Thr Val Leu Arg Gly Gly
85

<210> 87
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: NOV1c Primer 1

<400> 87
tcatcacatg acaacatgaa gctgt 25

<210> 88
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: NOV1c Primer 2

<400> 88
gaaaggccctc aaactctcca tctatg 26

<210> 89
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: NOV7a Primer 1

<400> 89
ccaatctctg atgccctgcg at 22

<210> 90
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: NOV7a Primer 2

<400> 90
aggtcagtgc cggagcctcc 20

<210> 91
<211> 19
<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag273b Forward

<400> 91
cggcttgacg atgcttcac 19

<210> 92

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag273b Probe

<400> 92
tgactttct gggcttacca atgctatttc aa 32

<210> 93

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag273b Reverse

<400> 93
gcacccatct caatatctgc aatattg 27

<210> 94

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag1094 Forward

<400> 94
atggactgga aaacctggaa 20

<210> 95

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag1094 Probe

<400> 95
tcctgcaaggc agataacaat tttatcaca 29

<210> 96
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag1094 Reverse

<400> 96
tgctaaaggc acttgggtca

20

<210> 97
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag2100 Forward

<400> 97
agatccctgg aacagaggat t

21

<210> 98
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag2100 Probe

<400> 98
tgtctgaagc caataaactt gcagca

26

<210> 99
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag2100 Reverse

<400> 99
ccttcatgtt cctttgggta a

21

<210> 100
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag217 Forward

<400> 100

atctgtgctg aggcatgttc ct 22

<210> 101
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag217 Probe

<400> 101
atcctcctcc ctccccggct ctc 23

<210> 102
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag217 Reverse

<400> 102
ctgcatggct ggtgtgatg 19

<210> 103
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag850 Forward

<400> 103
cctttcttct cttcctccctc aa 22

<210> 104
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag850 Probe

<400> 104
cacctggcga gtgctcctct ctg 23

<210> 105
<211> 19
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag850 Reverse

<400> 105

ggtggatggc gttgttagag

19

<210> 106

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag1469 Forward

<400> 106

cgtacgtctt ccatgatgag tt

22

<210> 107

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag1469 Probe

<400> 107

cgtggcctcg atgattaaga tccctt

26

<210> 108

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag1469 Reverse

<400> 108

aagtcaaggaa tgatggtaaa g

21

<210> 109

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag2976 Forward

<400> 109

acccaaatg gattccattt

20

<210> 110

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag2976 Probe

<400> 110
ccctcatgga tctgcataac cacaca 26

<210> 111

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag2976 Reverse

<400> 111
cttgtgtgtg catgcttg 20

<210> 112

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag760 Forward

<400> 112
caccatgaca acgacaccta ta 22

<210> 113

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag760 Probe

<400> 113
atatggcacc aacatcacat gcacg 25

<210> 114

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag760 Reverse

<400> 114
tgggttagaaa gtgtgtgtga aa 22

<210> 115
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag1537 Forward

<400> 115
aaggagctgg aagagaagaa ga 22

<210> 116
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag1537 Probe

<400> 116
atcagaaaact cagccctgga cacctg 26

<210> 117
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag1537 Reverse

<400> 117
gctgcgactt ggtcttgat 19

<210> 118
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag147 Forward

<400> 118
tgaagacagc acctccctat ca 22

<210> 119
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag147 Probe

<400> 119

cggctccgtg ctgtcaccca g

21

<210> 120
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag147 Reverse

<400> 120
aagaatcctc agcatcgcca ta

22

<210> 121
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag718 Forward

<400> 121
agaaggaatc tctggaaag ct

22

<210> 122
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag718 Probe

<400> 122
ccactggaga tgcttgtgtc tctacca

27

<210> 123
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag718 Reverse

<400> 123
gacagagcac tggcttagttc ac

22

<210> 124
<211> 22
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ag3681/Ag4085
Forward

<400> 124
gaatcatcca caagtcatcc at

22

<210> 125
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag3681/Ag4085
Probe

<400> 125
ctcaactccca tctcatgcct cccag

25

<210> 126
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Ag3681/Ag4085
Reverse

<400> 126
catgctgaat gatcgtgaca

20

<210> 127
<211> 1011
<212> PRT
<213> Homo sapiens

<400> 127
Ile Ser Asn Ser Ser Asp Thr Val Glu Cys Glu Cys Ser Glu Asn Trp
1 5 10 15

Lys Gly Glu Ala Cys Asp Ile Pro His Cys Thr Asp Asn Cys Gly Phe
20 25 30

Pro His Arg Gly Ile Cys Asn Ser Ser Asp Val Arg Gly Cys Ser Cys
35 40 45

Phe Ser Asp Trp Gln Gly Pro Gly Cys Ser Val Pro Val Pro Ala Asn
50 55 60

Gln Ser Phe Trp Thr Arg Glu Glu Tyr Ser Asn Leu Lys Leu Pro Arg
65 70 75 80

Ala Ser His Lys Ala Val Val Asn Gly Asn Ile Met Trp Val Val Gly
85 90 95

Gly Tyr Met Phe Asn His Ser Asp Tyr Asn Met Val Leu Ala Tyr Asp
 100 105 110

 Leu Ala Ser Arg Glu Trp Leu Pro Leu Asn Arg Ser Val Asn Asn Val
 115 120 125

 Val Val Arg Tyr Gly His Ser Leu Ala Leu Tyr Lys Asp Lys Ile Tyr
 130 135 140

 Met Tyr Gly Gly Lys Ile Asp Ser Thr Gly Asn Val Thr Asn Glu Leu
 145 150 155 160

 Arg Val Phe His Ile His Asn Glu Ser Trp Val Leu Leu Thr Pro Lys
 165 170 175

 Ala Lys Glu Gln Tyr Ala Val Val Gly His Ser Ala His Ile Val Thr
 180 185 190

 Leu Lys Asn Gly Arg Val Val Met Leu Val Ile Phe Gly His Cys Pro
 195 200 205

 Leu Tyr Gly Tyr Ile Ser Asn Val Gln Glu Tyr Asp Leu Asp Lys Asn
 210 215 220

 Thr Trp Ser Ile Leu His Thr Gln Gly Ala Leu Val Gln Gly Gly Tyr
 225 230 235 240

 Gly His Ser Ser Val Tyr Asp His Arg Thr Arg Ala Leu Tyr Val His
 245 250 255

 Gly Gly Tyr Lys Ala Phe Ser Ala Asn Lys Tyr Arg Leu Ala Asp Asp
 260 265 270

 Leu Tyr Arg Tyr Asp Val Asp Thr Gln Met Trp Thr Ile Leu Lys Asp
 275 280 285

 Ser Arg Phe Phe Arg Tyr Leu His Thr Ala Val Ile Val Ser Gly Thr
 290 295 300

 Met Leu Val Phe Gly Gly Asn Thr His Asn Asp Thr Ser Met Ser His
 305 310 315 320

 Gly Ala Lys Cys Phe Ser Ser Asp Phe Met Ala Tyr Asp Ile Ala Cys
 325 330 335

 Asp Arg Trp Ser Val Leu Pro Arg Pro Asp Leu His His Asp Val Asn
 340 345 350

 Arg Phe Gly His Ser Ala Val Leu His Asn Ser Thr Met Tyr Val Phe
 355 360 365

 Gly Gly Phe Asn Ser Leu Leu Leu Ser Asp Ile Leu Val Phe Thr Ser
 370 375 380

 Glu Gln Cys Asp Ala His Arg Ser Glu Ala Ala Cys Leu Ala Ala Gly
 385 390 395 400

Pro Gly Ile Arg Cys Val Trp Asn Thr Gly Ser Ser Gln Cys Ile Ser
 405 410 415

 Trp Ala Leu Ala Thr Asp Glu Gln Glu Glu Lys Leu Lys Ser Glu Cys
 420 425 430

 Phe Ser Lys Arg Thr Leu Asp His Asp Arg Cys Asp Gln His Thr Asp
 435 440 445

 Cys Tyr Ser Cys Thr Ala Asn Thr Asn Asp Cys His Trp Cys Asn Asp
 450 455 460

 His Cys Val Pro Arg Asn His Ser Cys Ser Glu Gly Gln Ile Ser Ile
 465 470 475 480

 Phe Arg Tyr Glu Asn Cys Pro Lys Asp Asn Pro Met Tyr Tyr Cys Asn
 485 490 495

 Lys Lys Thr Ser Cys Arg Ser Cys Ala Leu Asp Gln Asn Cys Gln Trp
 500 505 510

 Glu Pro Arg Asn Gln Glu Cys Ile Ala Leu Pro Glu Asn Ile Cys Gly
 515 520 525

 Ile Gly Trp His Leu Val Gly Asn Ser Cys Leu Lys Ile Thr Thr Ala
 530 535 540

 Lys Glu Asn Tyr Asp Asn Ala Lys Leu Phe Cys Arg Asn His Asn Ala
 545 550 555 560

 Leu Leu Ala Ser Leu Thr Thr Gln Lys Lys Val Glu Phe Val Leu Lys
 565 570 575

 Gln Leu Arg Ile Met Gln Ser Ser Gln Ser Met Ser Lys Leu Thr Leu
 580 585 590

 Thr Pro Trp Val Gly Leu Arg Lys Ile Asn Val Ser Tyr Trp Cys Trp
 595 600 605

 Glu Asp Met Ser Pro Phe Thr Asn Ser Leu Leu Gln Trp Met Pro Ser
 610 615 620

 Glu Pro Ser Asp Ala Gly Phe Cys Gly Ile Leu Ser Glu Pro Ser Thr
 625 630 635 640

 Arg Gly Leu Lys Ala Ala Thr Cys Ile Asn Pro Leu Asn Gly Ser Val
 645 650 655

 Cys Glu Arg Pro Ala Asn His Ser Ala Lys Gln Cys Arg Thr Pro Cys
 660 665 670

 Ala Leu Arg Thr Ala Cys Gly Asp Cys Thr Ser Gly Ser Ser Glu Cys
 675 680 685

 Met Trp Cys Ser Asn Met Lys Gln Cys Val Asp Ser Asn Ala Tyr Val
 690 695 700

Ala Ser Phe Pro Phe Gly Gln Cys Met Glu Trp Tyr Thr Met Ser Thr
 705 710 715 720

 Cys Pro Pro Glu Asn Cys Ser Gly Tyr Cys Thr Cys Ser His Cys Leu
 725 730 735

 Glu Gln Pro Gly Cys Gly Trp Cys Thr Asp Pro Ser Asn Thr Gly Lys
 740 745 750

 Gly Lys Cys Ile Glu Gly Ser Tyr Lys Gly Pro Val Lys Met Pro Ser
 755 760 765

 Gln Ala Pro Thr Gly Asn Phe Tyr Pro Gln Pro Leu Leu Asn Ser Ser
 770 775 780

 Met Cys Leu Glu Asp Ser Arg Tyr Asn Trp Ser Phe Ile His Cys Pro
 785 790 795 800

 Ala Cys Gln Cys Asn Gly His Ser Lys Cys Ile Asn Gln Ser Ile Cys
 805 810 815

 Glu Lys Cys Glu Asn Leu Thr Thr Gly Lys His Cys Glu Thr Cys Ile
 820 825 830

 Ser Gly Phe Tyr Gly Asp Pro Thr Asn Gly Gly Lys Cys Gln Pro Cys
 835 840 845

 Lys Cys Asn Gly His Ala Ser Leu Cys Asn Thr Asn Thr Gly Lys Cys
 850 855 860

 Phe Cys Thr Thr Lys Gly Val Lys Gly Asp Glu Cys Gln Leu Cys Glu
 865 870 875 880

 Val Glu Asn Arg Tyr Gln Gly Asn Pro Leu Arg Gly Thr Cys Tyr Tyr
 885 890 895

 Thr Leu Leu Ile Asp Tyr Gln Phe Thr Phe Ser Leu Ser Gln Glu Asp
 900 905 910

 Asp Arg Tyr Tyr Thr Ala Ile Asn Phe Val Ala Thr Pro Asp Glu Gln
 915 920 925

 Asn Arg Asp Leu Asp Met Phe Ile Asn Ala Ser Lys Asn Phe Asn Leu
 930 935 940

 Asn Ile Thr Trp Ala Ala Ser Phe Ser Ala Gly Thr Gln Ala Gly Glu
 945 950 955 960

 Glu Met Pro Val Val Ser Lys Thr Asn Ile Lys Glu Tyr Lys Asp Ser
 965 970 975

 Phe Ser Asn Glu Lys Phe Asp Phe Arg Asn His Pro Asn Ile Thr Phe
 980 985 990

 Phe Val Tyr Val Ser Asn Phe Thr Trp Pro Ile Lys Ile Gln Val Gln
 995 1000 1005

Thr Glu Gln
1010